

## SEQUENCE LISTING

<110> Wang, Tongtong  
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Watanabe, Yoshihiro  
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Peckham, David W.  
Fanger, Neil

<120> COMPOSITIONS AND METHODS FOR THE THERAPY  
AND DIAGNOSIS OF LUNG CANCER

<130> 210121.455C16

<140> US

<141> 2001-06-28

<160> 467

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 315

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 236, 241

<223> n = A,T,C or G

<400> 1

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cagctgccgt gagactcccg atgtcacagg cagtctgtgt ggttacagcg cccctcagtg 120
ttcatctcca gcagagacaa cggaggaggc tcccaccagg acggttctca ttatttatat 180
gttaatatgt ttgtaaactc atgtacagtt ttttttgggg gggaagcaat gggaanggta 240
naaattacaa atagaatcat ttgctgtaat ccttaaattgg caaacggtca ggccacgtga 300
aaaaaaaaaa aaaaaa                                     315
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<210> 2

<211> 380

<212> DNA

<213> Homo sapiens

<400> 2

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atntaggtt aagatthttgt ttacccttgt tactaaggag caaattagta ttaaagtata 60
atatatataa acaaatacaa aaagthttga gtggttcagc ttttttattt tttttaatgg 120
```

```

cataactttt aacaacactg ctctgtaatg ggttgaactg tggactcag actgagataa 180
ctgaaatgag tggatgtata gtgttattgc ataattatcc cactatgaag caaagggact 240
ggataaattc ccagtctaga ttattagcct ttgttaacca tcaagcacct agaagaagaa 300
ttattggaaa ttttgccttc tgtaactggc actttggggg gtgacttata ttttgccttt 360
gtaaaaaaaa aaaaaaaaaa

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<210> 3
<211> 346
<212> DNA
<213> Homo sapiens

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<220>
<221> misc_feature
<222> 316, 317, 318, 322, 323, 326, 329, 330, 331, 336, 337, 339,
340, 342, 343
<223> n = A,T,C or G

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<400> 3
ttgtaagtat acaatttttag aaaggattaa atgttattga tcatttttact gaatactgca 60
catcctcacc atacaccatc cactttccaa taacatttaa tcctttctaa aattgtaagt 120
atacaattgt actttctttg gattttcata acaaataac catagactgt taattttatt 180
gaagtttccct taatggaatg agtcattttt gtcttgtgct tttgaggtta cctttgcttt 240
gacttccaac aatttgatca tatagtgttg agctgtggaa atctttaagt ttattctata 300
gcaataattt ctatnnnnag annccngggn naaaannann annaaa

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<210> 4
<211> 372
<212> DNA
<213> Homo sapiens

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<220>
<221> misc_feature
<222> 297, 306, 332
<223> n = A,T,C or G

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<400> 4
actagtctca ttactccaga attatgctct tgtacctgtg tggctggggt tcttagtcgt 60
tggtttggtt tggttttttg aactgggatg taggggtggtt cacagttcta atgtaagcac 120
tctcttctcc aagttgtgct ttgtggggac aatcattctt tgaacattag agaggaaggc 180
agttcaagct gttgaaaaga ctattgctta tttttgtttt taaagaccta cttgacgtca 240
tgtggacagt gcacgtgcct tacgctacat cttgttttct aggaagaagg ggatgcnggg 300
aaggantggg tgctttgtga tggataaaaac gntaaataa cacaccttta cattttgaaa 360
aaaacaaaac aa

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<210> 5
<211> 698
<212> DNA
<213> Homo sapiens

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<220>
<221> misc_feature
<222> 8, 345, 422, 430, 433, 436, 438, 472, 481, 486, 515, 521,
536, 549, 553, 556, 557, 559, 568, 593, 597, 605, 611, 613,
616, 618, 620, 628, 630, 632, 634, 635, 639, 643, 647, 648,

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649, 652, 654, 658, 664, 690

<223> n = A,T,C or G

<400> 5

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actagtanga tagaaacact gtgtcccgag agtaaggaga gaagctacta ttgattagag 60
cctaaccagcag gttaactgca agaagaggcg ggatactttc agctttccat gtaactgtat 120
gcataaagcc aatgtagtcc agtttctaag atcatgttcc aagctaactg aatcccactt 180
caatacacac tcatgaactc ctgatggaac aataacaggc ccaagcctgt ggtatgatgt 240
gcacacttgc tagactcaga aaaaataacta ctctcataaa tgggtgggag tattttgggt 300
gacaacctac tttgcttggc tgagtgaagg aatgatattc atatnttcat ttattccatg 360
gacatttagt tagtgctttt tatataccag gcatgatgct gagtgcactc cttgtgtata 420
tntccaaatn ttingtnngt cgctgcacat atctgaaatc ctatattaag antttcccaa 480
natgangtcc ctgggttttc caccgacctt gatcngtcaa ngatctcacc tctgtntgtc 540
ctaaaacctn ctntctnnang gttagacnng acctctcttc tcccttcccg aanaatnaag 600
tgtgngaaga nancncncn cccccctn tcnncctng ccngctnnnc cnctgtngng 660
ggngnccgcc cccgcggggg gacccccccn ttttcccc 698
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<210> 6

<211> 740

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 82, 406, 426, 434, 462, 536, 551, 558, 563, 567, 582, 584,  
592, 638, 651, 660, 664, 673, 675, 697, 706, 711, 715, 716,  
717, 723, 724, 725, 733

<223> n = A,T,C or G

<400> 6

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actagtcaaa aatgctaaaa taatttgagg gaaaatattt ttttaagtagt gttatagttt 60
catgtttatc ttttattatg tnttgtgaag ttgtgtcttt tcactaatta cctatactat 120
gccaatattt ccttataatc atccataaca tttatactac atttgtaaga gaatatgcac 180
gtgaaactta acactttata aggtaaaaat gaggtttcca agatttaata atctgatcaa 240
gttcttggtt tttccaaata gaatggactt ggtctgttaa ggggctaagg gagaagaaga 300
agataagggt aaaagtgtgt aatgaccaa cattctaaaa gaaatgcaa aaaaaattta 360
ttttcaagcc ttogaactat ttaaggaaag caaatcatt tcctanatgc atatcatttg 420
tgagantttc tcantaatat cctgaatcat tcatctcagc tnaggcttca tgttgactcg 480
atatgtcatc tagggaaagt ctatttcatg gtccaaacct gttgccatag ttggtnaggc 540
tttcctttta ntgtgaanta ttnacangaa attttctctt tnanagttct tnatagggtt 600
aggggtgtgg gaaaagcttc taacaatctg tagtgtnng tgttatctgt ncagaaccan 660
aatnacggat cgnangaagg actgggtcta tttacangaa cgaatnatct ngttnnntgt 720
gtnnncaact ccngggagcc 740
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<210> 7

<211> 670

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 265, 268, 457, 470, 485, 546, 553, 566, 590, 596, 613, 624,  
639, 653, 659, 661

<223> n = A,T,C or G

<400> 7

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agcggccccc gctcgatggc cccgtggtgc tcagttagca gcggcccgtc gcgctacgtg 120
cttgggatgc aggagctgtt ccggggccac agcaagaccg cgagttcctg gcgcacagcg 180
ccaaggtgca ctcggtggcc tggagtgcg acgggcgtcg cctacctcgg ggtcttcgac 240
aagacgccac gtcttcttgc tgganaanga ccgttggtca aagaaaacaa ttatcgggga 300
catggggata gtgtggacca ctttgttggc atccaagtaa tcctgaccta tttgttacgg 360
cgtctggaga taaaaccatt cgcattctgg atgtgaggac tacaaaatgc attgccactg 420
tgaacactaa aggggagaac attaatatct gctggantcc tgatgggcan accattgctg 480
tagcnacaag gatgatgtgg tgactttatt gatgccaaga aaccccgttc caaagcaaaa 540
aaacanttcc aanttcgaag tcaccnaaat ctccctggaac aatgaacatn aatatnttct 600
tcctgacaat ggnccctggg tgtntcacat cctcagctnc cccaaaactg aancctgtnc 660
natccacccc
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<210> 8

<211> 689

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 253, 335, 410, 428, 448, 458, 466, 479, 480, 482, 483, 485,  
488, 491, 492, 495, 499, 500, 502, 503, 512, 516, 524, 525,  
526, 527, 530, 540, 546, 550, 581, 593, 594, 601, 606, 609,  
610, 620, 621, 622, 628, 641, 646, 656, 673

<223> n = A,T,C or G

<400> 8

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aaatgaagta ctggatttgg gaaaacctgg ttttattaga acatatggaa tgaaagccta 120
cacctagcat tgcctactta gccccctgaa ttaacagagc ccaattgaga caaacccctg 180
gcaacaggaa attcaaggga gaaaaagtaa gcaacttggg ctaggatgag ctgactccct 240
tagagcaaag ganagacagc ccccatcacc aaataccatt tttgcctggg gcttgtgcag 300
ctggcagtgt tcctgcccc aatgtggcacc ttatngtttt gatagcaact tcgttgaatt 360
ttcaccaact tattacttga aattataata tagcctgtcc gtttgcctgn tccaggctgt 420
gatataatnt cctagtgggt tgacttttna aataaatnag gtttantttt ctccccccnn 480
cnntnctncc nntcnctenn cnntcccccc cncctngtcc tccnnnttn gggggggccn 540
ccccncgggn ggacccccct ttgggtccctt agtggagggt natggcccct ggnnttatcc 600
nggcctann tttccccgtn nnaaatgntt cccctcccca ntccnccac ctcaanccgg 660
aagcctaagt ttntaccctg ggggtcccc
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<210> 9

<211> 674

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 602, 632, 639, 668

<223> n = A,T,C or G

<400> 9

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gtccactctc ctttgagtgt actgtcttac tgtgcactct gtttttcaac tttctagata 60
```



```
<210> 10
<211> 346
<212> DNA
<213> Homo sapiens
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|            |             |             |             |            |            |     |  |
|------------|-------------|-------------|-------------|------------|------------|-----|--|
| <400>      | 10          |             |             |            |            |     |  |
| actagtctgc | tgatagaaaag | cactatacat  | cctattgttt  | ctttctttcc | aaaatcagcc | 60  |  |
| ttctgtctgt | aacaaaaatg  | tactttatag  | agatggagga  | aaaggtctaa | tactacatag | 120 |  |
| ccttaagtgt | ttctgtcatt  | gttcaagtgt  | attttctgta  | acagaaacat | atttggaatg | 180 |  |
| tttttctttt | ccccttataa  | attgtaattc  | ctgaaatact  | gctgctttta | aaagtcccac | 240 |  |
| tgtcagatta | tattattctaa | caattggaata | ttgtaaatat  | actgtgtcta | cctctcaata | 300 |  |
| aaagggtact | tttctatttan | nnagnnqnnn  | qnnnnataaaa | anaaaaa    |            | 346 |  |

```
<210> 11
<211> 602
<212> DNA
<213> Homo sapiens
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```
<210> 12
<211> 685
<212> DNA
<213> Homo sapiens
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<220>  
 <221> misc\_feature  
 <222> 170, 279, 318, 321, 322, 422, 450, 453, 459, 467, 468, 470,  
 473, 475, 482, 485, 486, 491, 498, 503, 506, 509, 522, 526,  
 527, 528, 538, 542, 544, 551, 567, 568, 569, 574, 576, 582,  
 587, 588, 589, 590, 592, 593, 598, 599, 603, 605, 608  
 <223> n = A,T,C or G

<221> misc\_feature  
 <222> 633, 634, 635, 644, 646, 648, 651, 655, 660, 662, 663, 672,  
 674, 675, 682, 683  
 <223> n = A,T,C or G

<400> 12  
 actagtccctg tgaaagtaca actgaaggca gaaagtgtta ggatttttgca tctaattgttc 60  
 attatcatgg tattgatgga cctaagaaaa taaaaattag actaagcccc caaataagct 120  
 gcatgcattt gtaacatgat tagtagattt gaatatatag atgtagtatn ttgggtatct 180  
 aggtgtttta tcattatgta aaggaattaa agtaaaggac tttgtagttg tttttattaa 240  
 atatgcatat agtagagtgc aaaaatatag caaaaatana aactaaaggc agaaaagcat 300  
 tttagatatg ccttaatnta nnaactgtgc cagggtggccc tcggaataga tgccaggcag 360  
 agaccagtgc ctgggtgggtg cctccccttg tctgcccccc tgaagaactt cctcacgtg 420  
 angtagtgcc ctcgtaggtg tcacgtggan tantggganc aggccgnnnc gtnanaagaa 480  
 ancanngtga nagtttcncc gtngangcng aactgtccct gngccnnnac gctcccanaa 540  
 cntntccaat ngacaatcga gtttccnnnc tccngnaacc tngccgnnnn cnggcccnnc 600  
 cantntgnta accccgcgcc cggatcgctc tcnnttcgtt ctncncnaa ngggntttcn 660  
 cnnccgcgct cncnnccccg cnncc 685

<210> 13  
 <211> 694  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 503, 546, 599, 611, 636, 641, 643, 645, 656, 658, 662, 676,  
 679, 687  
 <223> n = A,T,C or G

<400> 13  
 cactagtcac tcattagcgt tttcaatagg gctcttaagt ccagtagatt acgggtagtc 60  
 agttgacgaa gatctgggtt acaagaacta attaaatgtt tcattgcatt tttgtaagaa 120  
 cagaataatt ttataaaatg tttgtagttt ataattgccg aaaataattt aaagacactt 180  
 tttctctgtg tgtgcaaagt tgtgtttgtg atccattttt tttttttttt taggacacct 240  
 gtttactagc tagcttttaca atatgccaaa aaaggatttc tccctgaccc catccgtggt 300  
 tcacctcttt ttccccccat gcttttttgc ctagttttata acaaaggaat gatgatgatt 360  
 taaaaagtat ttctgtatct tcagtatctt ggtcttccag aacctcttgg ttgggaaggg 420  
 gatcattttt tactggtcat ttcccttttg agtggtactac tttaacagat ggaaagaact 480  
 cattggccat ggaaacagcc gangtggttg gagccagcag tgcattggcac cgtccggcat 540  
 ctggentgat tgggtctggct gccgtcattg tcagcacagt gccatgggac atggggaana 600  
 ctgactgcac ngccaatggt tttcatgaag aatacngcat ncnngtgat cacgtnancc 660  
 angacgctat gggggncana gggccanttg cttc 694

<210> 14  
 <211> 679

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 29, 68, 83, 87, 94, 104, 117, 142, 145, 151, 187, 201, 211,  
226, 229, 239, 241, 245, 252, 255, 259, 303, 309, 359, 387,  
400, 441, 446, 461, 492, 504, 505, 512, 525, 527, 533, 574,  
592, 609, 610, 618, 620, 626, 627, 633, 639, 645, 654

<223> n = A,T,C or G

<400> 14

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cagccgcctg catctgtatc cagcgccang tcccgcctagt cccagctgcg cgcgcccccc 60
agtcccgna cccgttcggcc cangctnagt tagncctcac catnccggtc aaaggangca 120
ccaagtgcac caaataacctg cngtncggat ntaaattcat cttctggctt gccgggattg 180
ctgtccttgc cattggacta nggctccgat ncgactctca gaccanganc atcttcganc 240
naganactaa tnatnatntt tccagcttct acacaggagt ctatatcttg atcggatccg 300
gcncctctnt gatgctgggtg ggcttcctga gctgctgcgg ggctgtgcaa gagtcccant 360
gcatgctggg actgttcttc ggcttctctt tggtgataatn cgccattgaa atacctgcgg 420
ccatctgggg atattccact ncgatnatgt gattaaggaa ntccacggag ttttacaagg 480
acacgtacaa cnacctgaaa accnnggatg anccccaccg ggaancnctg aangccatcc 540
actatgcgtt gaactgcaat ggtttggctg gggnccttga acaattttaat cncatacatc 600
tggcccccann aaaggacntn ctcganncct tcnccgtgna attcngttct gatnccatca 660
cagaagtctc gaacaatcc                                     679
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<210> 15

<211> 695

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 105, 172, 176, 179, 189, 203, 212, 219, 221, 229, 231, 238,  
242, 261, 266, 270, 278, 285, 286, 298, 311, 324, 337, 350,  
363, 384, 391, 395, 405, 411, 424, 427, 443, 448, 453, 455,  
458, 463, 467, 470, 479, 482, 484, 493, 499, 505, 518

<223> n = A,T,C or G

<221> misc\_feature

<222> 520, 523, 531, 540, 584, 595, 597, 609, 611, 626, 628, 651,  
652, 657, 661, 665, 669, 672, 681, 683, 691, 693

<223> n = A,T,C or G

<400> 15

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actagtggat aaaggccagg gatgctgctc aacctcctac catgtacagg gacgtctccc 60
cattacaact acccaatccg aagtgtcaac tgtgtcagga ctaanaaacc ctggttttga 120
ttaaaaaagg gcctgaaaaa aggggagcca caaatctgtc tgcttcctca cnttantcnt 180
tggcaaatna gcattctgtc tcnttggctg cngcctcanc ncaaaaaanc ngaactcnat 240
cnggcccagg aatacatctc ncaatnaacn aaattganca aggcnnctgg aaatgccnga 300
tgggattatc ntccgcttgt tgancctcta agtttctntt ccttcattcn accctgccag 360
ccnagtctct ttagaaaaat gccngaattc naacnccggt tttctactc ngaatttaga 420
tctncanaaa ctctctggcc acnattcnaa ttanggnca cgnacanatn ccttccatna 480
ancncacccc acntttgana gccangacaa tgactgcntn aantgaaggc ntgaaggaan 540
aactttgaaa ggaaaaaaa ctttgtttcc ggcccccttc aacncttctg tgttnanac 600
```

```

tgccttctng naaccctgga agcccnnga cagtgttaca tgttggtcta nnaaacngac 660
ncttnaatnt cnatcttccc nanaacgatt ncnc 695

```

```

<210> 16
<211> 669
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 299, 354, 483, 555, 571, 573, 577, 642, 651, 662, 667
<223> n = A,T,C or G

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<400> 16
cgccgaagca gcagcgcagg ttgtccccgt ttccccctccc ccttcccttc tccggttgcc 60
ttcccgggcc ccttacactc cacagtcccg gtcccgccat gtcccagaaa caagaagaag 120
agaaccctgc ggaggagacc ggcgaggaga agcaggacac gcaggagaaa gaagggtattc 180
tgcctgagag agctgaagag gcaaagctaa aggccaaata cccaagccta ggacaaaagc 240
ctggaggctc cgacttcctc atgaagagac tccagaaagg gcaaaagtac tttgactcng 300
gagactacaa catggccaaa gccaacatga agaataagca gctgccaaagt gcangaccag 360
acaagaacct ggtgactggt gatcacatcc ccaccccaca ggatctgccc agagaaagtc 420
ctcgtctgtc accagcaagc ttgcgggtgg ccaagttgaa tgatgctgcc ggggctctgc 480
canatctgag acgtttccct ccctgccccca cccgggtcct gtgctggtc ctgcccttcc 540
tgcttttgca gccanggggtc aggaagtggc ncnggtngtg gctggaaagc aaaacccttt 600
cctgttggtg tcccacccat ggagcccctg ggcgagccc angaacttga ncctttttgt 660
tntcttnc 669

```

```

<210> 17
<211> 697
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 33, 48, 50, 55, 59, 60, 76, 77, 78, 90, 113, 118, 130, 135,
141, 143, 150, 156, 166, 167, 170, 172, 180, 181, 190, 192,
194, 199, 201, 209, 212, 224, 225, 226, 230, 233, 234, 236,
242, 244, 251, 253, 256, 268, 297, 305, 308, 311, 314
<223> n = A,T,C or G

```

```

<221> misc_feature
<222> 315, 317, 322, 324, 327, 333, 337, 343, 362, 364, 367, 368,
373, 384, 388, 394, 406, 411, 413, 423, 429, 438, 449, 450,
473, 476, 479, 489, 491, 494, 499, 505, 507, 508, 522, 523,
527, 530, 533, 535, 538, 539, 545, 548, 550, 552, 555
<223> n = A,T,C or G

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```

<221> misc_feature
<222> 562, 563, 566, 568, 572, 577, 578, 580, 581, 591, 594, 622,
628, 632, 638, 642, 644, 653, 658, 662, 663, 665, 669, 675,
680, 686, 689
<223> n = A,T,C or G

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```

<400> 17

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```

gcaagatatg gacaactaag tgagaaggta atnctctact gctctagntn ctcnnggenn 60
gacgcgctga ggagannnac gctggcccan ctgccggcca cacacgggga tcntggtnat 120
gcctgcccان gggancccca ncnctcgان cccatntcac acccgnnccn tncgccaان 180
ncctggetcn cncngccng nccagctcnc gncccccctcc gccnnnctcn ttnnctctc 240
cncnccctcc ncnacnacct cctaccncg gctccctccc cagccccccc ccgcaancct 300
ccacnacncc ntcnncnaga ancnccnctc gcnctcngcc ccnccccct gccccccgcc 360
cncnacnncg cgncccccg cgcncgcngc ctncccccct cccacnacag ncnacccgc 420
agnacgcnc tccgcccct gacgcccان cccgcgcgc tcaccttcac ggncnacng 480
ccccgctcnc ncnctgcnc gccgnccngg cgcgccgcc cnnccgngtn ccnccngng 540
ccccngcngn angcngtgcg cnnccangnc gngccggnnc ncacccctccg ncnccgcc 600
cgcccgctgg gggtcccgcc cncgcggntc antcccncc cntncgcca ctntccgntc 660
cnnnctcnc gctcngcgc cgcncncnc ccccccc 697

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<210> 18

<211> 670

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 234, 292, 329, 437, 458, 478, 487, 524, 542, 549, 550, 557, 576, 597, 603, 604, 646, 665

<223> n = A,T,C or G

<400> 18

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ctcgtgtgaa ggggtgcagta cctaagccgg agcggggtag aggcggggcg gcacccccctt 60
ctgacctcca gtgccgccgg cctcaagatc agacatggcc cagaacttga acgacttggc 120
gggacggctg cccgccgggc cccggggcat gggcacggcc ctgaagctgt tgctgggggc 180
cggcgccgtg gcctacgggtg tgcgcgaatc tgtgttcacc gtggaaggcg ggcncagagc 240
catcttcttc aatcggatcg gtggagtgc caggacacta tcctggggcg anggccttca 300
cttcaggatc cttgggtcca gtaccccanc atctatgaca ttcgggccag acctcgaaaa 360
aatctcctcc ctacaggctc caaagaccta cagatggtga atatctccct gcgagtgttg 420
tctcgaccaa tgctcangaa cttcctaaca tgttccancg cctaagggtt ggactacnaa 480
gaacgantgt tgccgtccat tgtcacgaag tgcacagaa tttnggtggc caagttcaat 540
gncctcacnn ctgatcnccc agcggggcca agttanccct ggttgatccc cgggganctg 600
acnnaaaagg gcccaaggact tcccctcatc ctggataatg tggcctcac aaagctcaac 660
tttanccacc 670

```

<210> 19

<211> 606

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 506

<223> n = A,T,C or G

<400> 19

```

actagtgccac acctcagctc ccaggccagt tctctgaatg tcgaggagtt ccaggatctc 60
tggcctcagt tgtccttgggt tattgatggg ggacaaattg gggatggcca gagccccgag 120
tgtgccttg gctcaactgt ggttgatttg tctgtgcccg gaaagtttgg catcattcgt 180
ccaggctgtg ccctggaaag tactacagcc atcctccaac agaagtacgg actgctcccc 240
tcacatgcgt cctacctgtg aaactctggg aagcaggaag gcccaagacc tgggtgctgga 300

```

```
tactatgtgt ctgtccactg acgactgtca aggcctcatt tgcagaggcc accggagcta 360
gggcactagc ctgactttta aggcagtgtg tctttctgag cactgtagac caagcccttg 420
gagctgctgg tttagccttg cacctgggga aaggatgtat ttatttgtat tttcatatat 480
cagccaaaag ctgaatggaa aagttnagaa cattcctagg tggccttatt ctaataagtt 540
tcttctgtct gttttgtttt tcaattgaaa agttattaaa taacagattt agaatctagt 600
gagacc 606
```

<210> 20

<211> 449

<212> DNA

<213> Homo sapiens

<400> 20

```
actagtaaac aacagcagca gaaacatcag tatcagcagc gtcgccagca ggagaatatg 60
cagcgccaga gccgaggaga acccccgctc cctgaggagg acctgtccaa actcttcaaa 120
ccaccacagc cgcctggccag gatggactcg ctgctcattg caggccagat aaacacttac 180
tgccagaaca tcaaggagtt cactgcccaa aacttaggca agctcttcat ggcccaggct 240
cttcaagaat acaacaacta agaaaaggaa gtttccagaa aagaagttaa catgaactct 300
tgaagtcaca ccagggcaac tcttggaaaga aatatatttg catattgaaa agcacagagg 360
atctctttag tgtcattgcc gatcttggct ataacagtgt ctttctagcc ataataaaat 420
aaaacaaaat cttgactgct tgctcaaaa 449
```

<210> 21

<211> 409

<212> DNA

<213> Homo sapiens

<400> 21

```
tatcaatcaa ctggtgaata attaaacaat gtgtggtgtg atcatacaaa gggtaccact 60
caatgataaa aggaacaagc tgcctatatg tggaacaaca tggatgcatt tcagaaactt 120
tatgttgagt gaaagaacaa acacggagaa catactatgt ggttctcttt atgtaacatt 180
acagaaataa aaacagaggc aaccaccttt gaggcagtat ggagtgagat agactggaaa 240
aaggaaggaa ggaaactcta cgctgatgga aatgtctgtg tcttcattgg gtggtagtta 300
tgtggggata tacatttgtc aaaatttatt gaactatata ctaaagaact ctgcatttta 360
ttgggatgta aataatacct caattaaaaa gacaaaaaaa aaaaaaaaaa 409
```

<210> 22

<211> 649

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 263, 353, 610, 635, 646

<223> n = A,T,C or G

<400> 22

```
acaattttca ttatcttaag cacattgtac atttctacag aacctgtgat tattctcgca 60
tgataaggat ggtacttgca tatggtgaat tactactgtt gacagtttcc gcagaaatcc 120
tatttcagtg gaccaacatt gtggcatggc agcaaatgcc aacattttgt ggaatagcag 180
caaatctaca agagaccctg gttggttttt cgttttgttt tctttgtttt ttcccccttc 240
tcctgaatca gcagggatgg aangagggtta gggaaagtat gaattactcc ttccagtagt 300
agctctgaag tgtcacattt aatatcagtt ttttttaaac atgattctag ttnaatgtag 360
aagagagaag aaagaggaag tgttcacttt ttttaatacac tgatttagaa atttgatgtc 420
```

```

ttatatcagt agttctgagg tattgatagc ttgctttatt tctgccttta cgttgacagt 480
gttgaagcag ggtgaataac taggggcata tatatttttt ttttttgtaa gctgtttcat 540
gatgttttct ttggaatttc cggataagtt caggaaaaca tctgcatgtt gttatctagt 600
ctgaagttcn tatccatctc attacaacaa aaacnccag aacggnnttg 649

```

```

<210> 23
<211> 669
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 642, 661
<223> n = A,T,C or G

```

```

<400> 23
actagtgccg tactggctga aatccctgca ggaccaggaa gagaaccagt tcagactttg 60
tactctcagt caccagctct ggaattagat aaattccttg aagatgtcag gaatgggac 120
tatcctctga cagccttttg gctgcctcgg cccagcagc cacagcagga ggaggtgaca 180
tcacctgtcg tgccccctc tgtcaagact ccgacacctg aaccagctga ggtggagact 240
cgcaaggtgg tgctgatgca gtgcaacatt gagtcggtgg aggagggagt caaacaccac 300
ctgacacttc tgctgaagtt ggaggacaaa ctgaaccggc acctgagctg tgacctgatg 360
ccaaatgaga atatccccga gttggcggct gagctggtgc agctgggctt cattagttag 420
gctgaccaga gccggttgac ttctctgcta gaagagactt gaacaagtgc aattttgcca 480
ggaacagtac cctcaactca gccgctgtca ccgtctcctc ttagagctca ctcgggccag 540
gccctgatct gcgctgtggc tgtcctggac gtgctgcacc ctctgtcctt cccccagtc 600
agtattacct gtgaagccct tccctccttt attattcagg anggctgggg gggtccttg 660
nttctaacc 669

```

```

<210> 24
<211> 442
<212> DNA
<213> Homo sapiens

```

```

<400> 24
actagtacca tcttgacaga ggatacatgc tcccaaaacg tttgttacca cacttaaaaa 60
tactgccat cattaagcat cagtttcaaa attatagcca ttcattgattt actttttcca 120
gatgactatc attattctag tcctttgaat ttgtaagggg aaaaaaaaca aaaacaaaaa 180
cttacgatgc acttttctcc agcacatcag atttcaaatt gaaaattaaa gacatgctat 240
ggtaatgcac ttgctagtac tacacacttt ggtacaacaa aaaacagagg caagaaacaa 300
cggaaagaga aaagccttcc tttgttggcc cttaaactga gtcaagatct gaaatgtaga 360
gatgatctct gacgatacct gtatgttctt attgtgtaaa taaaattgct ggtatgaaat 420
gacctaaaaa aaaaaaaga aa 442

```

```

<210> 25
<211> 656
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 330, 342, 418, 548, 579, 608
<223> n = A,T,C or G

```

```

<400> 25
tgcaagtacc acacactggt tgaattttgc acaaaaagtg actgtaggat caggtgatag 60
ccccggaatg tacagtgtct tgggtgcacca agatgccttc taaaggctga cataccttgg 120
accctaatagg ggcagagagt atagccctag cccagtgggtg acatgaccac tccctttggg 180
aggcctgagg tagaggggag tggatatgtg tttctcagtg gaagcagcac atgagtgggt 240
gacaggatgt tagataaagg ctctagttag ggtgtcattg tcatttgaga gactgacaca 300
ctcctagcag ctggtaaagg ggtgctggan gccatggagg anctctagaa acattagcat 360
gggctgatct gattacttcc tggcatcccg ctacttttta tgggaagtct tattagangg 420
atgggacagt tttccatata cttgctgtgg agctctggaa cactctctaa atttccctct 480
attaaaaatc actgccctaa ctacacttcc tccttgaagg aatagaaatg gaactttctc 540
tgacatantt cttggcatgg ggagccagcc acaaatgana atctgaacgt gtccagggtt 600
ctcctganac tcatctacat agaattgggt aaaccctccc ttggaataag gaaaaa 656

```

```

<210> 26
<211> 434
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 395
<223> n = A,T,C or G

```

```

<400> 26
actagttcag actgccacgc caaccccaga aaatacccca catgccagaa aagtgaagtc 60
ctagggtgtt ccatctatgt ttcaatctgt ccatctacca ggctctcgga taaaaacaaa 120
acaaaaaaac gctgccaggt tttagaagca gttctggtct caaaaccatc aggatcctgc 180
caccaggggt cttttgaaat agtaccacat gtaaaagggg atttggcttt cacttcatct 240
aataactgaa ttgtcaggct ttgattgata attgtagaaa taagtagcct tctgttgtgg 300
gaataagtta taatcagtat tcatctcttt gttttttgtc actcttttct ctctaattgt 360
gtcatttgta ctgtttgaaa aatatttctt ctatnaaatt aaactaacct gccttaaaaa 420
aaaaaaaaaa aaaa 434

```

```

<210> 27
<211> 654
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 505, 533, 563, 592, 613, 635, 638
<223> n = A,T,C or G

```

```

<400> 27
actagtccaa cacagtcaga aacattgttt tgaatcctct gtaaaccaag gcattaatct 60
taataaacca ggatccattt aggtaccact tgatataaaa aggatatcca taatgaatat 120
tttatactgc atcctttaca ttagccacta aatacgttat tgcttgatga agacctttca 180
cagaatccta tggattgcag catttcactt ggctacttca taccatgcc ttaaagaggg 240
gcagtttctc aaaagcagaa acatgccgcc agttctcaag ttttctcct aactccattt 300
gaatgtaagg gcagctggcc cccaatgtgg ggaggccga acattttctg aattcccatt 360
ttcttgttcg cggctaaatg acagtttctg tcattactta gattccgatc tttcccaaag 420
gtgttgattt acaaagaggc cagctaatag cagaaatcat gaccctgaaa gagagatgaa 480
attcaagctg tgagccaggc agganctcag tatggcaaag gtcttgagaa tcngccattt 540
ggtacaaaaa aaatttttaa gcntttatgt tataccatgg aaccatagaa anggcaaggg 600

```



aattgttaag aanaatttta agtgtccaga cccanaanga aaaaaaaaaa aaaa 654

<210> 28

<211> 670

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 101, 226, 274, 330, 385, 392, 397, 402, 452, 473, 476, 532, 534, 538, 550, 583, 595, 604, 613, 622, 643, 669

<223> n = A,T,C or G

<400> 28

```

cgtgtgcaca tactgggagg atttccacag ctgcacgggtc acagccctta cggattgcc 60
ggaaggggag aaagatatgt gggataaact gagaaaagaa nccaaaaacc tcaacatcca 120
aggcagctta ttcgaactct gcggcagcgg caacggggcg gcgggggtccc tgctcccggc 180
gttcccgggt ctctgggtgt ctctctcggc agcttttagcg acctgncctt ccttctgagc 240
gtggggccag ctccccccgc ggcgcccaacc cacnctcact ccatgctccc ggaaatcgag 300
aggaagatca ttagttcttt ggggacgttn gtgattctct gtgatgctga aaaacactca 360
tatagggaat gtgggaaatc ctganctctt tnttatntcg tntgatttct tgtgttttat 420
ttgccaaaat gttaccaatc agtgaccaac cnagcacagc caaaaatcgg acntcngctt 480
tagtccgtct tcacacacag aataagaaaa cggcaaacc accccacttt tnantttnat 540
tattactaan ttttttctgt tgggcaaaa aatctcagga acngccctgg ggccnccgta 600
ctanagttaa ccnagctagt tncatgaaaa atgatgggct ccnccctcaat gggaaagcca 660
agaaaaagnc 670

```

<210> 29

<211> 551

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 336, 474, 504, 511, 522, 523, 524, 540, 547

<223> n = A,T,C or G

<400> 29

```

actagtcttc cacagcctgt gaatccccct agacctttca agcatagtga gcggagaaga 60
agatctcagc gtttagccac cttacccatg cctgatgatt ctgtagaaaa ggtttcttct 120
ccctctccag cactgatgg gaaagtattc tccatcagtt ctcaaaatca gcaagaatct 180
tcagtaccag agtgccctga tgttgacat ttgccacttg agaagctggg accctgtctc 240
cctcttgact taagtcgtgg ttcagaagtt acagcaccgg tagcctcaga ttcctcttac 300
cgtaatgaat gtcccagggc agaaaaagag gatacncaga tgcttccaaa tccttcttcc 360
aaagcaatag ctgatgggaa gaggagctcc agcagcagca ggaatatcga aaacagaaaa 420
aaaagtgaat ttgggaagac aaaagctcaa cagcatttgg taaggagaaa aganaagatg 480
aggaaggaag agagaagaga gacnaagatc nctacggacc gnncggaag aagaagaagn 540
aaaaanaaaa a 551

```

<210> 30

<211> 684

<212> DNA

<213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 545, 570, 606, 657, 684  
 <223> n = A,T,C or G

<400> 30  
 actagttcta tctggaaaaa gcccgggttg gaagaagctg tggagagtgc gtgtgcaatg 60  
 cgagactcat ttcttggaag catccctggc aaaaatgcag ctgagtacaa ggttatcact 120  
 gtgatagaac ctggactgct ttttgagata atagagatgc tgcagtctga agagacttcc 180  
 agcacctctc agttgaatga attaataatgatg gcttctgagt caactttact ggctcaggaa 240  
 ccacgagaga tgactgcaga tgtaatcgag cttaaaggga aattcctcat caacttagaa 300  
 ggtggtgata ttctgtgaaga gtcttcctat aaagtaattg tcatgccgac tacgaaagaa 360  
 aaatgcccc gttgttgga gtatacagcg ggagtcctta gatacactgt gtccctcgatg 420  
 tgcagaagtt gtcagtggga aaatagtatt aacagctcac tcgagcaaga accctcctga 480  
 cagtactggg ctagaagttt ggatggatta tttacaatat aggaaagaaa gccaagaatt 540  
 aggtnatgag tggatgagta aatggtggan gatggggaat tcaaatacaga attatggaag 600  
 aagtnttcc tgttactata gaaaggaatt atgtttatatt acatgcagaa aatatanatg 660  
 tgtggtgtgt accgtggatg gaan 684

<210> 31  
 <211> 654  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 326, 582, 651  
 <223> n = A,T,C or G

<400> 31  
 gcgcagaaaa ggaaccaata tttcagaaac aagcttaata ggaacagctg cctgtacatc 60  
 aacatcttct cagaatgacc cagaagttat catcgtggga gctggcgtgc ttggctctgc 120  
 ttggcagct gtgctttcca gagatggaag aaagggtgaca gtcattgaga gagacttaaa 180  
 agagcctgac agaatagttg gagaattcct gcagccgggt gggtatcatg ttctcaaaga 240  
 ccttggctctt ggagatacag tggaaaggtct tgatgccag gttgtaaatg gttacatgat 300  
 tcatgatcag ggaaagcaaa tcagangttc agattcctta ccctctgtca gaaaacaatc 360  
 aagtgcagag tggaagagct ttccatcacg gaagattcat catgagtctc cggaaagcag 420  
 ctatggcaga gcccaatgca aagtttattg aagggtgtgt gttacagtta ttagaggaag 480  
 atgatgttgt gatgggagtt cagtacaagg ataaagagac tgggagatat caaggaactc 540  
 catgctccac tgactgttgt tgcagatggg cttttctcca anttcaggaa aagcctgggtc 600  
 tcaataaagt ttctgtatca ctcatttggt tggcttctta tgaagaatgc nccc 654

<210> 32  
 <211> 673  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 376, 545, 627  
 <223> n = A,T,C or G

<400> 32  
 actagtgaag aaaaagaaat tctgatacgg gacaaaaatg ctcttcaaaa catcattctt 60

```

tatcacctga caccaggagt ttctattgga aaaggatttg aacctgggtg tactaacatt 120
ttaaagacca cacaaggaag caaaatcttt ctgaaagaag taaatgatac acttctggtg 180
aatgaattga aatcaaaaga atctgacatc atgacaacaa atggtgtaat tcatgttgta 240
gataaactcc tctatccagc agacacacct gttggaaatg atcaactgct ggaaatactt 300
aataaattaa tcaaatacat ccaaattaag tttgttcgtg gtagcacctt caaagaaatc 360
cccgtagctg tctatnagcc aattattaaa aaatacacca aaatcattga tgggagtgcc 420
tgtgggaaat aactgaaaaa gagaccgaga agaacgaatc attacaggtc ctgaaataaa 480
atacctagga tttctactgg aggtggagaa acagaagaac tctgaagaaa ttgttacaag 540
aagangtccc aaggtcacca aattcattga aggtgggtgat ggtctttatt tgaagatgaa 600
gaaattaaaa gacgcttcag ggagacnccc catgaaggaa ttgccagcca caaaaaaatt 660
cagggattag aaa 673

```

```

<210> 33
<211> 673
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 325, 419, 452, 532, 538, 542, 571, 600, 616, 651, 653, 672
<223> n = A,T,C or G

```

```

<400> 33
actagttatt tactttcctc cgcttcagaa ggtttttcag actgagagcc taagcatact 60
ggatctgttg tttcttttgg gtctcacctc atcagtgtgc atagtggcag aaattataaa 120
gaaggttgaa aggagcaggg aaaagatcca gaagcatggt agttcgacat catcatcttt 180
tcttgaagta tgatgcatat tgcattatit tatttgcaaa ctaggaattg cagtctgagg 240
atcatttaga agggcaagtt caagaggata tgaagatttg agaacttttt aactattcat 300
tgactaaaaa tgaacattaa tgttnaagac ttaagacttt aacctgctgg cagtcccaaa 360
tgaaattatg caactttgat atcatattcc ttgatttaaa ttgggctttt gtgattgant 420
gaaactttat aaagcatatg gtcagttatt tnattaaaaa ggcaaaacct gaaccacctt 480
ctgcacttaa agaagtctaa cagtacaaat acctatctat cttagatgga tntatttntt 540
tntattttta aatattgtac tatttatggg nggtggggct ttcttactaa tacacaaatn 600
aatttatcat ttcaanggca ttctatttgg gtttagaagt tgattccaag nantgcatat 660
ttcgctactg tnt 673

```

```

<210> 34
<211> 684
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 414, 472, 480, 490, 503, 507, 508, 513, 523, 574, 575, 598,
659, 662, 675
<223> n = A,T,C or G

```

```

<400> 34
actagtttat tcaagaaaag aacttactga ttcctctgtt cctaaaagcaa gagtggcagg 60
tgatcagggc tgggtgtagca tccggttcct ttagtgagc taactgcatt tgtcactgat 120
gaccaaggag gaaatcacta agacatttga gaagcagtg tatgaacgtt cttggacaag 180
ccacagttct gagccttaac cctgtagttt gcacacaaga acgagctcca cctccccttc 240
ttcaggagga atctgtgcgg atagattggc tggacttttc aatggttctg ggttgcaagt 300
gggcactggt atggctgggt atggagcgga cagccccagg aatcagagcc tcagcccggc 360

```

```

tgccctggttg gaaggtacag gtgttcagca ccttcggaaa aagggcataa agtngtgggg 420
gacaattctc agtccaagaa gaatgcattg accatttgctg gctatttgct tncctagtan 480
gaattggatn catttttgac cangatnntt ctncatgctt ttnttgcaat gaaatcaaat 540
cccgcaattat ctacaagtgg tatgaagtcc tgcnnccccc agagaggctg ttcaggcnat 600
gtcttccaag ggcagggtgg gttacaccat tttacctccc ctctcccccc agattatgna 660
cncagaagga atttntttcc tccc                                     684

```

```

<210> 35
<211> 614
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 17, 20, 152, 223, 267, 287, 304, 306, 316, 319, 321, 355,
365, 382, 391, 407, 419, 428, 434, 464, 467, 477, 480, 495,
499, 505, 515, 516, 522, 524, 527, 542, 547, 549, 567, 572,
576, 578
<223> n = A,T,C or G

```

```

<400> 35
actagtccaa cgcgttngcn aatattcccc tggtagccta cttccttacc cccgaatatt 60
ggtaagatcg agcaatggct tcaggacatg gggtctcttc tcctgtgatc attcaagtgc 120
tactgtcatg aagactggct tgtctcagtg tntcaacctc accagggctg tctcttggtc 180
cacacctcgc tccctgttag tgccgtatga cagcccccat canatgacct tggccaagtc 240
acggtttctc tgtggtcaat gttggtnngc tgattggtgg aaagtanggt ggaccaaaagg 300
aagncncgtg agcagncanc nccagttctg caccagcagc gcctccgtcc tactnggggtg 360
ttcngtcttc tcctggccct gngtgggcta nggctgatt cggaanattg cctttgcang 420
gaaggganga taantgggat ctaccaattg attctggcaa aacnatntct aagattnttn 480
tgctttatgt ggganacana tctanctctc atttnntgct gnanatnaca cctactcgt 540
gntcgancnc gtcttcgatt ttcgganaca cnccantnaa tactggcggt ctgttggtta 600
aaaaaaaaaa aaaa                                     614

```

```

<210> 36
<211> 686
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 222, 224, 237, 264, 285, 548, 551, 628, 643, 645, 665, 674
<223> n = A,T,C or G

```

```

<400> 36
gtggctggcc cggttctccg cttctcccca tcccctactt tcctccctcc ctccctttcc 60
ctccctcgct gactgttgct tgctggtcgc agactccctg acccctccct caccctcccc 120
taacctcggt gccaccgat tgcccttctt ttctgttgct ccagcccgag cctagtgtca 180
gggcgggggc ctggagcagc ccgaggcact gcagcagaag ananaaaaga cacgacnaac 240
ctcagctcgc cagtcgggtc gctngcttcc cgccgcatgg caatnagaca gacgccgctc 300
acctgctctg ggcacacgcg acccggtggtt gatttgccct tcagtggcat cacccttatg 360
ggtattttctt aatcagcgct tgcaaagatg gttaacctat gctacgccag ggagatacag 420
gagactggat tggaacattt ttggggctcta aaggtctggt tggggtgcaa cactgaataa 480
ggatgccacc aaagcagcta cagcagctgc agatttcaca gcccaagtgt gggatgctgt 540
ctcagganat naattgataa cctgggtcat aacacattgt caagaatgtg gatttcccca 600

```

```

ggatattatt atttgtttac cgggggganag gataactggt tcnctatatt taattgaaca 660
aactnaaaca aaanctaagg aaatcc 686

```

```

<210> 37
<211> 681
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 7, 10, 11, 19, 25, 32, 46, 53, 77, 93, 101, 103, 109, 115,
123, 128, 139, 157, 175, 180, 192, 193, 194, 212, 218, 226,
227, 233, 240, 241, 259, 260, 267, 289, 296, 297, 298, 312,
313, 314, 320, 325, 330, 337, 345, 346, 352, 353, 356
<223> n = A,T,C or G

```

```

<221> misc_feature
<222> 382, 385, 400, 427, 481, 484, 485, 491, 505, 515, 533, 542,
544, 554, 557, 560, 561, 564, 575, 583, 589, 595, 607, 619,
628, 634, 641, 645, 658, 670
<223> n = A,T,C or G

```

```

<400> 37
gagacanacn naacgtcang agaanaaaag angcatggaa cacaanccag gncgatggc 60
caccttccca ccagcancca gcgcccccca gngcccgga ngnccggang accangactc 120
cancctgnat caatctganc tctattcctg gcccatncct acctcggagg tggangccgn 180
aaaggctgca cnnncagaga agctgctgcc ancaccancc gcccnnccc tgnccgggctn 240
nataggaaac tgggtgaccnn gctgcanaat tcatacagga gcacgcgang ggcacnnnct 300
cacactgagt tnnngatgan gctnaccan ggacctnccc cagcnnattg annacnggac 360
tgccggaggaa ggaagacccc gnaacggatc ctggccggcn tgccaccccc ccacccttag 420
gattatnccc cttgactgag tctctgagg gctaccgaa cccgcctcca ttcctacca 480
natnntgctc natcgggact gacangctgg ggatnggagg ggctatcccc cancatcccc 540
tnanaccaac agcnaangan natnggggct ccccnngggtc gngcacaacnc tctcncaccc 600
cggcgcnggc cttcgggtgt tctctcctc aacnaattcc naaanggcgg gccccccngt 660
ggactcctcn ttgttcctc c 681

```

```

<210> 38
<211> 687
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 3, 30, 132, 151, 203, 226, 228, 233, 252, 264, 279, 306,
308, 320, 340, 347, 380, 407, 429, 437, 440, 445, 448, 491,
559, 567, 586, 589, 593, 596, 603, 605, 606, 609, 626, 639,
655, 674, 682
<223> n = A,T,C or G

```

```

<400> 38
canaaaaaaa aaaacatggc cgaaaccagn aagctgcgcg atggcgccac ggccccctctt 60
ctcccgccct gtgtccggaa ggtttccctc cgaggcgccc cggtctccgc aagcggagga 120
gagggcgga cntgccggg ccggagctca naggccctgg ggccgctctg ctctcccgcc 180
atcgcaaggg cggcgctaac ctnaggcctc cccgcaaagg tccccnangc gngggcggcg 240

```

```

gggggctgtg anaaccgcaa aaanaacgct gggcgcgcn ggaacccgtc ccccccgcg 300
aaggananac ttccacagan gcagcgtttc cacagccan agccacnttt ctaggggtgat 360
gcaccccagt aagttcctgn cggggaagct caccgctgtc aaaaaanctc ttgctccac 420
cggcgcacna agggggangan ggcangangc tgccgcccgc acagggtcatc tgatcacgtc 480
gcccgcccta ntctgctttt gtgaatctcc actttgttca accccacccg ccgttctctc 540
ctccttgccg cttcctctna ccttaanaac cagcttcctc taccnctng tanttntct 600
gcncnngtng aaattaattc ggtcnccgg aacctcttnc ctgtggcaac tgctnaaaga 660
aactgctgtt ctgnttactg cngtccc 687

```

```

<210> 39
<211> 695
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 300, 401, 423, 429, 431, 437, 443, 448, 454, 466, 492, 515,
523, 524, 536, 538, 541, 552, 561, 566, 581, 583, 619, 635,
636, 641, 649, 661, 694
<223> n = A,T,C or G

```

```

<400> 39
actagtctgg cctacaatag tgtgattcat gtaggacttc tttcatcaat tcaaaacccc 60
tagaaaaacg tatacagatt atataagtag ggataagatt tctaacattt ctgggctctc 120
tgacccttgc gctagactgt ggaaaggag tattattata gtatacaaca ctgctgttgc 180
cttattagtt ataacatgat aggtgctgaa ttgtgattca caatttaaaa acactgtaat 240
ccaaactttt ttttttaact gtagatcatg catgtgaatg ttaatgttaa tttgttcaan 300
gttgttatgg gtagaaaaaa ccacatgcct taaaatttta aaaagcaggg cccaaactta 360
ttagtttaaa attaggggta tgtttccagt ttgttattaa ntggttatag ctctgtttag 420
aanaaatcna ngaacangat ttngaaantt aagntgacat tatttnccag tgacttgta 480
atttgaaatc anacacggca ccttccggtt tggtntctatt ggnnttgaa tccaancngg 540
ntccaaatct tnttggaac ngtcnttta acttttttac nanatcttat ttttttattt 600
tggaatggcc ctatttaang ttaaaagggg ggggnnccac naccattcnt gaataaaact 660
naatatatat ccttgggtccc ccaaaattta aggng 695

```

```

<210> 40
<211> 674
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 403, 428, 432, 507, 530, 543, 580, 583, 591, 604, 608, 621,
624, 626, 639, 672
<223> n = A,T,C or G

```

```

<400> 40
actagtagtc agttgggagt ggttgctata ccttgacttc atttatatga atttccactt 60
tattaaataa tagaaaagaa aatcccgggtg cttgcagtag agttatagga cattctatgc 120
ttacagaaaa tatagccatg attgaaatca aatagtaaag gctgttcttg ctttttatct 180
tcttagctca tcttaataaa gtagtacact tgggatgcag tgcgtctgaa gtgctaata 240
gttgtaacaa tagcacaat cgaacttagg atgtgtttct tctcttctgt gtttcgattt 300
tgatcaattc ttttaatttg ggaacctata atacagtttt cctattcttg gagataaaaa 360
ttaaatggat cactgatatt taagtcattc tgcttctcat ctnaatattc catattctgt 420

```

```

attagganaa antacctccc agcacagccc cctctcaaac cccacccaaa accaagcatt 480
tggaatgagt ctccctttatt tccgaantgt ggatgggtata acccatatcn ctccaatttc 540
tgnttggggtt ggggtattaat ttgaactgtg catgaaaagn ggnaatcttt nctttgggtc 600
aaanttttccc ggttaatttg nctngncaaa tccaatttnc ttttaagggtg tctttataaa 660
atttgctatt cngg 674

```

```

<210> 41
<211> 657
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 243, 247, 251, 261, 267, 272, 298, 312, 315, 421, 432, 434,
501, 524, 569, 594, 607, 650
<223> n = A,T,C or G

```

```

<400> 41
gaaacatgca agtaccacac actgtttgaa ttttgcacaa aaagtgactg tagggatcag 60
gtgatagccc cggaatgtac agtgtcttgg tgcaccaaga tgccttctaa aggctgacat 120
accttgggac cctaattgggg cagagagtat agccctagcc cagtgggtgac atgaccactc 180
cctttgggag gctgaagtta aagggaatgg tatgtgtttt ctcatggaag cagcacatga 240
atnggtnaca ngatgttaaa ntaaggntct antttgggtg tcttgtcatt tgaaaaantg 300
acacactcct ancanctggg aaaggggtgc tggagccat ggaagaactc taaaaacatt 360
agcatgggct gatctgatta ctccctggca tcccgcacac ttttatggga agtcttatta 420
naaggatggg ananttttcc atatccttgc tgttgggaact ctggaacact ctctaaattt 480
ccctctatta aaaatcactg nccttactac acttcctcct tganggaata gaaatggacc 540
tttctctgac ttagttcttg gcatggganc cagcccaaat taaaatctga cttntccggt 600
ttctccngaa ctcacctact tgaattggta aaacctcctt tgggaattagn aaaaacc 657

```

```

<210> 42
<211> 389
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 179, 317, 320
<223> n = A,T,C or G

```

```

<400> 42
actagtgctg aggaatgtaa acaagtttgc tgggccttgc gagacttcac caggttggtt 60
cgatagctca cactcctgca ctgtgcctgt caccacaggaa tgtctttttt aattagaaga 120
caggaagaaa acaaaaacca gactgtgtcc cacaatcaga aacctccgtt gtggcagang 180
ggccttcacc gccaccaggg tgtcccgcga gacagggaga gactccagcc ttctgaggcc 240
atcctgaaga attcctgttt ggggggttgtg aaggaaaatc acccgattt aaaaagatgc 300
tgttgcctgc ccgcgtngtn ggggaaggac tggtttcctg gtgaatttct taaaagaaaa 360
atattttaag ttaagaaaaa aaaaaaaaaa 389

```

```

<210> 43
<211> 279
<212> DNA
<213> Homo sapiens

```

```

<400> 43
actagtgcac agctcctggt cttgagatgt cttctcggtta aggagatggg ccttttggag 60
gtaaaggata aaatgaatga gttctgtcat gattcactat tctagaactt gcatgacctt 120
tactgtgtta gctctttgaa tgttcttgaa atttttagact ttctttgtaa acaaataata 180
tgtccttatac attgtataaa agctgttatg tgcaacagtg tggagatcct tgtctgattt 240
aataaaatac ttaaacactg aaaaaaaaaa aaaaaaaaaa 279

```

```

<210> 44
<211> 449
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 245, 256, 264, 266, 273, 281, 323, 325, 337, 393
<223> n = A,T,C or G

```

```

<400> 44
actagtagca tcttttctac aacgttaaaa ttgcagaagt agcttatcat taaaaaaca 60
caacaacaac aataacaata aatcctaagt gtaaatcagt tattctaccc cctaccaagg 120
atatcagcct gttttttccc ttttttctcc tgggaataat tgtgggcttc ttcccaaatt 180
tctacagcct ctttcctctt ctcatgcttg agcttccctg ttgcaacgca tgcgttgtgc 240
aagantgggc tgtttngctt ggantncggt ccnagtggaa ncatgctttc ccttggttact 300
gttggaagaa actcaaacct tcnancccta ggtgttncca ttttgtcaag tcatcactgt 360
atttttgtac tggcattaac aaaaaaagaa atnaaatatt gttccattaa actttaataa 420
aactttaaaa gggaaaaaaa aaaaaaaaaa 449

```

```

<210> 45
<211> 559
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 263
<223> n = A,T,C or G

```

```

<400> 45
actagtgtgg gggaatcacg gacacttaaa gtcaatctgc gaaataattc ttttattaca 60
cactcactga agtttttgag tcccagagag ccatttctatg tcaaacattc caagtactct 120
ttgagagccc agcattacat caacatgccc gtgcagttca aaccgaagtc cgcaggcaaa 180
tttgaagctt tgcttgtcat tcaaacagat gaaggcaaga gtattgctat tcgactaatt 240
ggtgaagctc ttggaaaaaa ttnactagaa tactttttgt gttaagttaa ttacataagt 300
tgtattttgt taactttatc tttctacact acaattatgc ttttgtatat atattttgta 360
tgatggatat ctataattgt agattttgtt tttaacaagct aatactgaag actcgactga 420
aatattatgt atctagccca tagtattgta cttaactttt acagggtgaa aaaaaaattc 480
tgtgtttgca ttgattatga tattctgaat aaatatggga atatatttta atgtgggtaa 540
aaaaaaaaaa aaaaaggaa 559

```

```

<210> 46
<211> 731
<212> DNA
<213> Homo sapiens

```



<220>  
 <221> misc\_feature  
 <222> 270, 467, 477, 502, 635, 660, 671, 688, 695, 697, 725  
 <223> n = A,T,C or G

<400> 46  
 actagttcta gtaccatggc tgtcatagat gcaaccatta tattccattt agtttcttcc 60  
 tcaggttccc taacaattgt ttgaaactga atatatatgt ttatgtatgt gtgtgtgttc 120  
 actgtcatgt atatggtgta tatgggatgt gtgcagtttt cagttatata tatattcata 180  
 tatacatatg catatatatg tataatatac atatatacat gcatacactt gtataaatata 240  
 catatatata cacatatatg cacacatatn atcactgagt tccaaagtga gtcttttattt 300  
 ggggcaattg tattctctcc ctctgtctgc tcaactgggcc tttgcaagac atagcaattg 360  
 cttgatttcc tttggataag agtcttatct tcggcactct tgactctagc cttaacttta 420  
 gatttctatt ccagaatacc tctcatatct atcttaaaac ctaaganggg taaagangtc 480  
 ataagattgt agtatgaaa antttgctta gttaaattat atctcaggaa actcattcat 540  
 ctacaaatta aattgtaaaa tgatggtttg ttgtatctga aaaaatgttt agaacaagaa 600  
 atgtaactgg gtacctgtta tatcaaagaa cctcnattta ttaagtctcc tcatagccan 660  
 atccttatat ngccctctct gacctgannt aatananact tgaataatga atagttaatt 720  
 taggnntggg c 731

<210> 47  
 <211> 640  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 5, 28, 106, 153, 158, 173, 176, 182, 189, 205, 210, 214,  
 225, 226, 229, 237, 260, 263, 269, 277, 281, 282, 322, 337,  
 338, 354, 365, 428, 441, 443, 456, 467, 476, 484, 503, 508,  
 554, 567, 575, 579, 588, 601, 606, 609, 611, 621, 636  
 <223> n = A,T,C or G

<400> 47  
 tgcnggccgg tttggccctt ctttgtanga cactttcatt cgccctgaaa ttttcccgat 60  
 cgtaataaac tctcagggtc cctgcctgca cagggttttt tcttantttg ttgcctaaca 120  
 gtacaccaaa tgtgacatcc tttcaccaat atngatttct tcataccaca tcntcnatgg 180  
 anacgactnc aacaattttt tgatnaccn aaanactggg ggctnnaana agtacantct 240  
 ggagcagcat ggacctgtcn gcnactaang gaacaanagt nntgaacatt tacacaacct 300  
 ttggtatgtc ttactgaaa anagaaacat gcttctnncc ctagaccacg aggncaaccg 360  
 caganattgc caatgccaag tccgagcggg tagatcagg aatacattcc atggatgcat 420  
 tacatacnth gtccccgaaa nanaagatgc cctaanggct tcttcanact ggccngaaa 480  
 acanctacac ctggtgcttg ganaacanac tctttggaag atcatctggc acaagttccc 540  
 ccagtggggt tttnctttgg cactanctt accanactna ttcggaancc attctttggc 600  
 ntggenttnt ntggggacca ntcttctcac aactgnaccc 640

<210> 48  
 <211> 257  
 <212> DNA  
 <213> Homo sapiens

<400> 48  
 actagtatat gaaaatgtaa atatcacttg tgtactcaaa caaaagttgg tcttaagctt 60  
 ccaccttgag cagccttgga aacctaacct gcctctttta gcataatcac attttctaaa 120

```

tgattttctt  tgttcctgaa  aaagtgattt  gtattagttt  tacatttggt  ttttggaaga  180
ttatatttgt  atatgtatca  tcataaaaata  tttaaataaa  aagtatcttt  agagtgaaaa  240
aaaaaaaaaa  aaaaaaa                                257

```

```

<210> 49
<211> 652
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 410, 428, 496, 571, 647
<223> n = A,T,C or G

```

```

<400> 49
actagttcag  atgagtggct  gctgaagggg  ccccttggtc  attttcatta  taacccaatt  60
tccacttatt  tgaactctta  agtcataaat  gtataatgac  ttatgaatta  gcacagttaa  120
gttgacacta  gaaactgccc  atttctgtat  tacactatca  aataggaaac  attggaaaga  180
tggggaaaaa  aatcttattt  taaaatggct  tagaaagttt  tcagattact  ttgaaaattc  240
taaacttctt  tctgtttcca  aaacttgaaa  atatgtagat  ggactcatgc  attaagactg  300
ttttcaaagc  ttctctcaca  tttttaaaag  gtgattttcc  ttttaatatata  catatttatt  360
ttctttaaag  cagctatatc  ccaacccatg  actttggaga  tatacctatn  aaaccaatat  420
aacagcangg  ttattgaagc  agctttctca  aatgttgctt  cagatgtgca  agttgcaaat  480
tttattgtat  ttgtanaata  caatttttgt  tttaaactgt  atttcaatct  atttctccaa  540
gatgcttttc  atatagagtg  aaatatccca  ngataactgc  ttctgtgtcg  tcgcatttga  600
cgcataactg  cacaaatgaa  cagtgtatac  ctcttggttg  tgcattnacc  cc                                652

```

```

<210> 50
<211> 650
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 237, 270, 311, 443, 454, 488, 520, 535, 539, 556, 567, 594,
603, 634
<223> n = A,T,C or G

```

```

<400> 50
ttgcgctttg  atttttttag  ggcttggtgc  ctgtttcact  tatagggtct  agaatgcttg  60
tgttgagtaa  aaaggagatg  cccaatattc  aaagctgcta  aatgttctct  ttgccataaa  120
gactccgtgt  aactgtgtga  acacttgga  tttttctcct  ctgtcccag  gtcgtcgtct  180
gctttctttt  ttgggttctt  tctagaagat  tgagaaatgc  atatgacagg  ctgagancac  240
ctcccaaac  acacaagctc  tcagccacan  gcagcttctc  cacagcccca  gcttcgcaca  300
ggctctgga  nggctgcctg  ggggaggcag  acatgggagt  gccaaggtgg  ccagatgggt  360
ccaggactac  aatgtcttta  tttttaactg  tttgccactg  ctgccctcac  ccctgcccgg  420
ctctggagta  ccgtctgccc  canacaagtg  ggantgaaat  gggggtgagg  gggaacactg  480
attcccantt  agggggtgcc  taactgaaca  gtagggatan  aagggtgtga  cctgngaant  540
gcttttataa  attatnttcc  ttgttanatt  tattttttta  tttaatctct  gttnaactgc  600
ccngggaaaa  ggggaaaaaa  aaaaaaaaaa  tctnttttaa  cacatgaaca                                650

```

```

<210> 51
<211> 545
<212> DNA

```

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 66, 159, 195, 205, 214, 243, 278, 298, 306, 337, 366, 375, 382, 405, 446, 477, 492, 495, 503, 507, 508, 521, 537

<223> n = A,T,C or G

<400> 51

```
tggcgtgcaa ccagggtagc tgaagtttgg gtctgggact ggagattggc cattaggcct 60
cctganattc cagctccctt ccaccaagcc cagtcttgct acgtggcaca gggcaaacct 120
gactcccttt gggcctcagt ttccccctccc cttcatgana tgaaaagaat actacttttt 180
cttgttggtc taacnttgct ggacncaaag tgtngtcatt attgttgat tgggtgatgt 240
gtncaaaact gcagaagctc actgcctatg agaggaanta agagagatag tggatganag 300
ggacanaagg agtcattatt tggatatagat ccaccntcc caacctttct ctcctcagtc 360
cctgncctc atgtntctgg tntgggtgagt cctttgtgcc accanccatc atgctttgca 420
ttgtgccat cctgggaagg ggggtgnatcg tctcacaact tgttgatc gtttganatg 480
catgctttct tnatnaaaca aanaaannaa tgtttgacag ngttttaaata aaaaaanaaa 540
caaaa
```

<210> 52

<211> 678

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 98, 119, 121, 131, 136, 139, 140, 142, 143, 163, 168, 172, 176, 184, 189, 190, 191, 200, 201, 205, 207, 221, 223, 229, 230, 237, 240, 241, 255, 264, 266, 267, 276, 280, 288, 289, 291, 297, 301, 306, 308, 314, 315, 326, 332, 335, 337

<223> n = A,T,C or G

<221> misc\_feature

<222> 339, 341, 343, 344, 345, 347, 350, 355, 356, 358, 362, 363, 372, 379, 395, 397, 398, 400, 403, 412, 414, 421, 423, 431, 435, 438, 439, 450, 457, 463, 467, 471, 474, 480, 483, 484, 487, 490, 491, 492, 493, 499, 500, 504, 508, 518, 536

<223> n = A,T,C or G

<221> misc\_feature

<222> 538, 549, 551, 552, 554, 556, 557, 562, 563, 567, 571, 572, 576, 579, 590, 592, 595, 598, 606, 609, 613, 620, 622, 624, 626, 631, 634, 638, 641, 647, 654, 660, 661, 674

<223> n = A,T,C or G

<400> 52

```
actagtagaa gaactttgcc gcttttgtgc ctctcacagg cgcctaaagt cattgccatg 60
ggaggaagac gatttggggg gggagggggg gggggcangg tccgtggggc tttccctant 120
ntatctccat ntccantgnn cnntgtcgcc tcttccctcg tcn cattnga anttantccc 180
tggccccnn nccctctccn nccnccct cccccctccg ncnccctccn cttttntan 240
ncttccccat ctcctcccc cctnanngtc ccaacnccgn cagcaatnnc ncacttnctc 300
nctccnccncc tccnccgtt cttctnttct cnactntnnc ncnntnccn tgccnntnaa 360
annctctccc cntgcaanc gattctctcc ctcnccnnan cnttccactc cntncttctc 420
```

```

nncgctcct nttctcnnc ccacctctcn ccttcgnccc cantacnctc nccncccttn 480
cgnntcnttn nntcctcnn accncccncc tcccttcncc cctctttctcc ccggtntntc 540
tctctcccncc nncnncncc cncnccntcc nngcgncnt ttcgccccn cncnccntt 600
ccttctcncc cantccatcn cntntnccat nctnccctncc nctcacnccc gctnccccn 660
ntctctttca cacngtcc 678

```

```

<210> 53
<211> 502
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 139, 146, 215, 217, 257, 263, 289, 386, 420, 452, 457, 461,
466, 482, 486
<223> n = A,T,C or G

```

```

<400> 53
tgaagatcct ggtgtcgcca tgggcccgcg ccccgcccgt tgttaccggt attgtaagaa 60
caagccgtac ccaaagtctc gcttctgccg aggtgtccct gatgccaaaa ttgcatttt 120
tgacctgggg cggaaaaaang caaaantgga tgagtctccg ctttgtggcc acatggtgtc 180
agatcaatat gagcagctgt cctctgaagc cctgnanget gcccgaattt gtgccaataa 240
gtacatggta aaaagtngtg gcnaagatgc ttccatatcc ggggtgcgnt ccaccccttc 300
cacgtcatcc gcatcaacaa gatgttgtcc tgtgctgggg ctgacaggct cccaacaggc 360
atgcgaagtg cctttggaaa acccanggca ctgtggccag gggttcacatt gggccaattn 420
atcatgttca tccgcaccaa ctgcagaaca angaactgt naattnaagc cctgcccagg 480
gncaanttca aatttcccgg cc 502

```

```

<210> 54
<211> 494
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 431, 442, 445
<223> n = A,T,C or G

```

```

<400> 54
actagtccaa gaaaaatatg cttaatgtat attacaaagg ctttgtatat gttaacctgt 60
tttaatgccaa aaagtttgct ttgtccacaa ttctcttaag acctcttcag aaagggattt 120
gtttgcctta atgaatactg ttgggaaaaa acacagtata atgagtgaag agggcagaag 180
caagaaattt ctacatctta gcgactccaa gaagaatgag tatccacatt tagatggcac 240
attatgagga ctttaatctt tccttaaaca caataatgtt ttcttttttc ttttattcac 300
atgatttcta agtatatttt tcatgcagga cagtttttca accttgatgt acagtgactg 360
tgttaaattt ttctttcagt ggcaacctct ataactttta aaatatggtg agcatcttgt 420
ctgttttgaa ngggatatga cnatnaatct atcagatggg aaatcctgtt tccaagttag 480
aaaaaaaaaa aaaa 494

```

```

<210> 55
<211> 606
<212> DNA
<213> Homo sapiens

```

<220>  
 <221> misc\_feature  
 <222> 375, 395, 511, 542, 559, 569, 578, 581  
 <223> n = A,T,C or G

<400> 55  
 actagtaaaa agcagcattg ccaaataatc cctaattttc cactaaaaat ataatgaaat 60  
 gatgttaagc tttttgaaaa gttaggtta aacctactgt tgtagatta atgtatttgt 120  
 tgcttccctt tatctggaat gtggcattag cttttttatt ttaacctctt ttaattctta 180  
 ttcaattcca tgacttaagg ttggagagct aaacactggg atttttggat aacagactga 240  
 cagttttgca taattataat cggcattgta catagaaagg atatggctac cttttgttaa 300  
 atctgcactt tctaaatatc aaaaaaggga aatgaagtat aaatcaattt ttgtataatc 360  
 tgtttgaaac atgantttta ttgtcttaat attanggctt tgcccttttc tgtagtctc 420  
 ttgggatcct gtgtaaaact gttctcatta aacaccaaac agttaagtcc attctctggt 480  
 actagctaca aattccgttt catattctac ntaacaattt aaattaactg aaatatttct 540  
 anatggtcta cttctgtcnt ataaaaacna aacttgantt nccaaaaaaa aaaaaaaaaa 600  
 aaaaaa 606

<210> 56  
 <211> 183  
 <212> DNA  
 <213> Homo sapiens

<400> 56  
 actagtatat ttaaacttac aggcttattt gtaatgtaaa ccaccatttt aatgtactgt 60  
 aattaacatg gttataatac gtacaatcct tccctcatcc catcacacaa ctttttttgt 120  
 gtgtgataaa ctgatttttg tttgcaataa aaccttgaaa aataaaaaaaaa aaaaaaaaaa 180  
 aaa 183

<210> 57  
 <211> 622  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 358, 368, 412, 414, 425, 430, 453, 455, 469, 475, 495, 499,  
 529, 540, 564, 575, 590  
 <223> n = A,T,C or G

<400> 57  
 actagtcaact actgtcttct ccttgtagct aatcaatcaa tattcttccc ttgcctgtgg 60  
 gcagtggaga gtgctgctgg gtgtacgctg cacctgcca ctgagttggg gaaagaggat 120  
 aatcagtgag cactgttctg ctcagagctc ctgatctacc ccaccccta ggatccagga 180  
 ctgggtcaaa gctgcatgaa accaggccct ggcagcaacc tgggaatggc tggaggtggg 240  
 agagaacctg acttctcttt cctctccct cctccaacat tactggaact ctatcctggt 300  
 agggatcttc tgagcttggt tccctgctgg gtgggacaga agacaaaagga gaagggangg 360  
 tctacaanaa gcagcccttc tttgtcctct ggggttaatg agcttgacct ananttcatg 420  
 gaganaccan aagcctctga tttttaattt cctnnaaatg tttgaagtnt atatntacat 480  
 atatataatt ctttnaatnt ttgagtcttt gatatgtctt aaaatccant ccctctgccn 540  
 gaaacctgaa ttaaaaccat gaanaaaaaat gtttncctta aagatgttan taattaattg 600  
 aaacttgaaa aaaaaaaaaa aa 622

<210> 58

<211> 433  
 <212> DNA  
 <213> Homo sapiens

<400> 58  
 gaacaaattc tgaattggta tgtaccgtca aaagacttga agaaatttca tgattttgca 60  
 gtgtggaagc gttgaaaatt gaaagttact gcttttccac ttgctcatat agtaaagga 120  
 tcctttcagc tgccagtgtt gaataatgta tcatccagag tgatgttata tgtgacagtc 180  
 accagcttta agctgaacca ttttatgaat accaaataaa tagacctctt gtactgaaaa 240  
 catatttgtg actttaatcg tgctgcttgg atagaaatat ttttactggg tcttctgaat 300  
 tgacagtaaa cctgtccatt atgaatggcc tactgttcta ttatttggtt tgacttgaat 360  
 ttatccacca aagacttcat ttgtgtatca tcaataaagt tgtatgtttc aactgaaaaa 420  
 aaaaaaaaaa aaa 433

<210> 59  
 <211> 649  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 22, 190, 217, 430, 433, 484, 544, 550, 577, 583, 594  
 <223> n = A,T,C or G

<400> 59  
 actagttatt atctgacttt cnggttataa tcatttctaag gagtgtgaag tagcctctgg 60  
 tgtcatttgg atttgcatth ctctgatgag tgatgctatc aagcaccttt gctgggtgctg 120  
 ttggccatat gtgtatgttc cctggagaag tgtctgtgct gagccttggc ccacttttta 180  
 attaggcgtt tgccttttta ttactgagtt gtaaganttc tttatatatt ctggattcta 240  
 gacccttatac agatacatgg tttgcaaata ttttctccca ttctgtgggt tgtgttttca 300  
 ctttatcgat aatgtcctta gacatataat aaatttgtat tttaaaagtg acttgatttg 360  
 ggctgtgcaa ggtgggctca cgcttgtaat ccagcactt tgggagactg aggtgggtgg 420  
 atcatatgan gangctagga gtctcgaggtc agcctggcca gcatagcgaa aacttgtctc 480  
 tacnaaaaat acaaaaatta gtcaggcatg gtgggtgcacg tctgtaatac cagcttctca 540  
 ggangctgan gcacaaggat cacttgaacc ccagaangaa gangttgcag tganctgaag 600  
 atcatgccag ggcaacaaaa atgagaactt gtttaaaaaa aaaaaaaaaa 649

<210> 60  
 <211> 423  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 209, 222, 277, 389, 398  
 <223> n = A,T,C or G

<400> 60  
 actagttcag gccttccagt tcaactgacaa acatggggaa gtgtgcccag ctggctggaa 60  
 acctggcagt gataccatca agcctgatgt ccaaaagagc aaagaatatt tctccaagca 120  
 gaagtgcagc ctgggctgtt ttagtgccag gctgcggtgg gcagccatga gaacaaaacc 180  
 tcttctgtat tttttttttc cattagtana acacaagact cngattcagc cgaattgtgg 240  
 tgtcttacaa ggcagggctt tcctacaggg ggtgganaaa acagcctttc ttcctttggg 300  
 aggaatggcc tgagttggcg ttgtgggcag gctactgggt tgtatgatgt attagtagag 360

```

caaccatta atcttttgta gtttgatna aacttganct gagaccttaa acaaaaaaaaa 420
aaa 423

```

```

<210> 61
<211> 423
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 195, 285, 295, 329, 335, 340, 347, 367, 382, 383, 391, 396,
418
<223> n = A,T,C or G

```

```

<400> 61
cgggactgga atgtaaagtg aagttcggag ctctgagcac gggctcttcc cgccgggtcc 60
tccctcccca gacccagag ggagaggccc accccgccc gccccgccc agccctgct 120
caggtctgag tatggctggg agtcgggggc cacaggcctc tagctgtgct gctcaagaag 180
actggatcag ggtanctaca agtggccggg ccttgccctt gggattctac cctgttccta 240
atttgggtgt ggggtgcggg gtccctggcc cctttttcca cactncctcc ctcngacag 300
caacctccct tggggcaatt gggcctggnt ctcncccg ttttgcnaacc ctttgttgg 360
ttaaggncct taaaaatgtt annttttccc ntgccnggt taaaaaagga aaaaactnaa 420
aaa 423

```

```

<210> 62
<211> 683
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 218, 291, 305, 411, 416, 441, 443, 453, 522, 523, 536, 542,
547, 566, 588, 592, 595, 603, 621, 628, 630, 632, 644, 645,
648, 655, 660, 672, 674, 676, 677, 683
<223> n = A,T,C or G

```

```

<400> 62
gctggagagg ggtacggact ttcttggagt tgtcccaggt tggaatgaga ctgaactcaa 60
gaagagaccc taagagactg gggaatggtt cctgccttca ggaaagtga agacgcttag 120
gctgtcaaca cttaaaggaa gtccccttga agcccagagt ggacagacta gacccattga 180
tggggccact ggccatggtc cgtggacaag acattccngt gggccatggc acaccggggg 240
ggatcaaaat gtgtacttgt ggggtctcgc cccttgccaa aaccaaacca ntcccactcc 300
tgtcnttggg ctttcttccc attccctcct ccccaaatgc acttcccctc ctccctctgc 360
ccctcctgtg tttttggaat tctgtttccc tcaaaattgt taatttttta nttttngacc 420
atgaacttat gtttgggtgc nangttcccc ttnccaatgc atactaatat attaatgggt 480
atattttttt gaaatatatt ttaatgaact tggaaaaaat tnntggaatt tccttncttc 540
cntttntttt ggggggggtg ggggngtggg ttaaaatttt tttggaancc cnatnggaaa 600
ttnttacttg gggccccctt naaaaaantn anttccaatt cttnnatngc ccctnttccn 660
ctaaaaaaaa ananannaaa aan 683

```

```

<210> 63
<211> 731
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 237, 249, 263, 288, 312, 317, 323, 326, 337, 352, 362, 370,
377, 400, 411, 414, 434, 436, 446, 457, 473, 486, 497, 498,
502, 512, 531, 546, 554, 563, 565, 566, 588, 597, 608, 611,
613, 615, 627, 632, 640, 641, 644, 654, 660, 663, 665
<223> n = A,T,C or G

```

```

<221> misc_feature
<222> 671, 678, 692, 697, 698, 699, 704, 705, 712, 714, 717, 718,
719, 723, 725, 730, 731
<223> n = A,T,C or G

```

```

<400> 63
actagtcata aaggggtgtgc gcgtcttcga cgtggcggtc ttggcgccac tgctgcgaga 60
cccgcccttg gacctcaagg tcatccactt ggtgcgtgat ccccgcgagg tggcgagttc 120
acggatccgc tcgcgccacg gcctcatccg tgagagccta cagggtggtgc gcagccgaga 180
ccgcgagctc accgcatgcc cttcttgagg gccgcgggcc acaagcttgg cgcccanaaa 240
gaaggcgtn ggggcccgc aantaccacg ctctgggcgc tatggaangt cctcttgcaa 300
taatattggt tnaaaanctg canaanagcc cctgcanccc cctgaactgg gntgcagggc 360
cncttacctn gtttggnctg gggtacaaag aacctgtttn ggaaaaccct nccnaaaacc 420
ttccgggaaa attntncaaa ttttntttgg ggaattnttg ggtaaaccct ccnaaaatgg 480
gaaacntttt tgccctnnaa antaaaccat tnggttccgg gggccccccc ncaaaaccct 540
ttttnttttt ttntgcccc cantnncccc ccggggcccc ttttttngg ggaaaanccc 600
ccccctncc nanantttta aaagggnggg anaatttttn nttncccccc gggncccccn 660
ggngntaaaa nggtttcncc ccccgagggg gnggggnnnc ctcnnaaacc cntntcnna 720
ccncttttn n 731

```

```

<210> 64
<211> 313
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 240
<223> n = A,T,C or G

```

```

<400> 64
actagttgtg caaaccacga ctgaagaaag acgaaaagtg ggaaataact tgcaacgtct 60
gttagagatg gttgctacac atgttgggtc tgtagagaaa catcttgagg agcagattgc 120
taaagttgat agagaatatg aagaatgcat gtcagaagat ctctcggaat atattaaaga 180
gattagagat aagtatgaga agaaagctac tctaattaag tcttctgaag aatgaagatn 240
aaatgttgat catgtatata tatccatagt gaataaaatt gtctcagtaa agttgtaaaa 300
aaaaaaaaaa aaa 313

```

```

<210> 65
<211> 420
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature

```



<222> 400, 402, 403, 404, 405, 406, 409, 411, 412, 414, 415, 416

<223> n = A,T,C or G

<400> 65

```
actagttccc tggcaggcaa gggcttccaa ctgaggcagt gcatgtgtgg cagagagagg 60
caggaagctg gcagtggcag cttctgtgtc tagggagggg tgtggctccc tccttccttg 120
tctgggaggt tggagggaag aatctaggcc ttagcttgcc ctctgccac cttccccctt 180
gtagatactg ccttaacact cctcctcttc tcagctgtgg ctgccacca agccaggttt 240
ctccgtgtct actaatttat ttccaggaaa ggtgtgtgga agacatgagc cgtgtataat 300
atttgtttta acattttcat tgcaagtatt gaccatcatc cttggttgtg tatcgttgta 360
acacaaatta atgatattaa aaagcatcca aacaaagccn annnnnaana nnnnnngaaa 420
```

<210> 66

<211> 676

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 328, 454, 505, 555, 586, 612, 636, 641

<223> n = A,T,C or G

<400> 66

```
actagtttcc tatgatcatt aaactcattc tcagggttaa gaaaggaatg taaatttctg 60
cctcaatttg tacttcatca ataagttttt gaagagtgca gatttttagt cagggtcttaa 120
aaataaaactc acaaatctgg atgcatttct aaattctgca aatgtttcct ggggtgactt 180
aacaaggaat aatcccacaa tatacctagc tacctaatac atggagctgg ggctcaaccc 240
actgttttta aggatttgcg cttacttgtg gctgaggaaa aataagtagt tccgagggaa 300
gtagttttta aatgtgagct tatagatngg aaacagaata tcaacttaat tatggaaatt 360
gttagaaacc tgttctcttg ttatctgaat cttgattgca attactattg tactggatag 420
actccagccc attgcaaagt ctcatatc ttanctgtgt agttgaattc cttggaaatt 480
ctttttaaga aaaaattgga gtttnaaaga aataaacccc tttgttaaat gaagcttggc 540
tttttgggtga aaaanaatca tcccgcaggg cttattgttt aaaaanggaa ttttaagcct 600
ccctggaaaa anttgtaaat taaatgggga aaatgntggg naaaaattat ccgttagggg 660
ttaaagggaa aactta 676
```

<210> 67

<211> 620

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 419, 493, 519, 568, 605, 610

<223> n = A,T,C or G

<400> 67

```
caccattaaa gctgcttacc aagaacttcc ccagcatttt gacttccttg tttgatagct 60
gaattgtgag cagggtgatag aagagccttt ctagttgaac atacagataa tttgctgaat 120
acattccatt taatgaaggg gttacatctg ttacgaagct actaagaagg agcaagagca 180
taggggaaaa aaatctgatc agaacgcac aaactcacat gtgccccctc tactacaaac 240
agattgtagt gctgtggtgg tttattccgt tgtgcagaac ttgcaagctg agtcactaaa 300
cccaaagaga ggaaattata ggtagttaa acattgtaat ccaggaact aagttaatt 360
```

```

cacttttgaa gtgttttggt ttttattttt gggttgctctg atttactttg ggggaaaang 420
ctaaaaaaaa agggatatca atctctaatt cagtgccac taaaagttgt ccctaaaaag 480
tctttactgg aanttatggg actttttaag ctccaggntt tttggctctc caaatataacc 540
ttgcatgggc cccttaaaat tgttgaangg cattcctgcc tctaagtttg gggaaaattc 600
ccccnttttn aaaatttgga                                     620

```

```

<210> 68
<211> 551
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 286, 464, 480, 501, 502, 518, 528, 533, 536, 537, 538, 539,
540, 541, 543, 544, 545, 547, 548, 549
<223> n = A,T,C or G

```

```

<400> 68
actagtagct ggtacataat cactgaggag ctatttctta acatgctttt atagaccatg 60
ctaagtctag accagtatit aagggtctaat ctacacacct cttagctgta agagtctggc 120
ttagaacaga cctctctgtg caataacttg tggccactgg aaatccctgg gccggcattt 180
gtattggggg tgcaatgact cccaagggcc aaaagagtta aaggcacgac tgggatttct 240
tctgagactg tggtgaaact ccttccaagg ctgagggggg cagtangtgc tctgggaggg 300
actcggcacc actttgatat tcaacaagcc acttgaagcc caattataaa attgttattt 360
tacagctgat ggaactcaat ttgaaccttc aaaactttgt tagtttatcc tatttatattg 420
ttaaaccctaa ttacatttgt ctagcattgg atttgggttc tgtngcatat gtttttttcn 480
cctatgtgct cccctccccc nnatcttaat ttaaaccnca attttgcnat tcncnnnnnn 540
nannnnanna a                                     551

```

```

<210> 69
<211> 396
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 235, 310, 323, 381
<223> n = A,T,C or G

```

```

<400> 69
cagaaatgga aagcagagtt ttcatttctg tttataaacg tctccaaaca aaaatggaaa 60
gcagagtttt cattaaatcc ttttaccttt ttttttctt ggtaatcccc tcaataaaca 120
gtatgtggga tattgaatgt taaagggata ttttttcta ttatttttat aattgtacaa 180
aattaagcaa atgttaaaaag ttttatatgc tttattaatg ttttcaaaag gtatnatata 240
tgtgatacat tttttaagct tcagttgctt gtcttctggt actttctgtt atgggctttt 300
ggggagccan aaaccaatct acnatctctt tttgtttgcc aggacatgca ataaaattta 360
aaaaataaat aaaaactatt nagaaattga aaaaaa                                     396

```

```

<210> 70
<211> 536
<212> DNA
<213> Homo sapiens

```

```

<220>

```

<221> misc\_feature  
 <222> 388, 446, 455  
 <223> n = A,T,C or G

<400> 70

```
actagtgc aaagcaaatat aaacatcgaa aaggcggtcc tcacgttagc tgaagatata 60
cttcgaaaga cccctgtaaa agagcccaac agtgaaaatg tagatatcag cagtggagga 120
ggcgtgacag gctggaagag caaatgctgc tgagcattct cctgttccat cagtggccat 180
ccactacccc gttttctctt cttgctgcaa aataaaccac tctgtccatt tttaaactcta 240
aacagatatt tttgtttctc atcttaacta tccaagccac ctatttttatt tgttctttca 300
tctgtgactg cttgctgact ttatcataat tttcttcaaa caaaaaaatg tatagaaaaa 360
tcatgtctgt gacttcattt ttaaagtnta cttgctcagc tcaactgcat ttcagttggt 420
ttatagtcga gttcttatca acattnaaac ctatngcaat cttttcaaat ctattctgca 480
aattgtataa gaataaaaag tagaatttaa caattaaaaa aaaaaaaaaa aaaaaa 536
```

<210> 71

<211> 865

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 22, 35, 39, 56, 131, 138, 146, 183, 194, 197, 238, 269, 277,  
 282, 297, 316, 331, 336, 340, 341, 346, 349, 370, 376, 381,  
 382, 392, 396, 397, 401, 433, 444, 445, 454, 455, 469, 472,  
 477, 480, 482, 489, 497, 499, 511, 522, 526, 527

<223> n = A,T,C or G

<221> misc\_feature

<222> 545, 553, 556, 567, 574, 580, 610, 613, 634, 638, 639, 663,  
 672, 689, 693, 694, 701, 704, 713, 723, 729, 732, 743, 744,  
 749, 761, 765, 767, 769, 772, 774, 780, 783, 788, 792, 803,  
 810, 824, 840, 848

<223> n = A,T,C or G

<400> 71

```
gacaaagcgt taggagaaga anagaggcag ggaanactnc ccaggcacga tggccncctt 60
cccaccagca accagcgccc cccaccagcc cccaggcccc gacgacgaag actccatcct 120
ggattaatct nacctctntc gcctgnccca ttcctacctc ggaggtggag gccggaaaagg 180
tcncaccaag aganaantctg ctgccaacac caaccgcccc agccctggcg ggcacganag 240
gaaactgggtg accaatctgc agaattctna gaggaanaag cnaggggccc cgcgctnaga 300
cagagctgga tatgangcca gaccatggac nctacncccn ncaatncana cgggactgcg 360
gaagatggan gaccncgcac nngatcaggc cngctnncca nccccccacc cctatgaatt 420
attcccgcgtg aangaatctc tgannggctt ccannaaagc gcctccccnc cnaacgnaan 480
tncaacatng ggattanang ctgggaactg naaggggcaa ancctnnaat atccccagaa 540
acaantcttc ccnaanaaac tggggcncct catnggtggn aactaaaccg 600
cacgccaagn aantataaaa ggggggcccc tcncggnng accccctttt gtcccttaat 660
ganggttata cnccttgctg accatgggtnc ccnnttctgt ntgnatgttt ccnctccctt 720
ccnctatnt cnagccgaac tcnnatttnc ccgggggtgc natcnantng tncnctttt 780
ttngttgncc cngccctttc cngcgggaacn cgtttccccc ttantaacgg caccgggggn 840
aagggtgntt ggccccctcc ctccc 865
```

<210> 72

<211> 560

<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> 83, 173, 183, 186, 209, 211, 215, 255, 321, 322, 323, 335,  
344, 357, 361, 368, 394, 412, 415, 442, 455, 469, 472, 475,  
487, 513, 522, 528, 531, 534, 546  
<223> n = A,T,C or G

<400> 72  
cctggacttg tcttgggtcc agaacctgac gacccggcga cggcgacgtc tcttttgact 60  
aaaagacagt gtccagtgtc ccngcctagg agtctacggg gaccgcctcc cgcgccgcca 120  
ccatgcccaa cttctctggc aactggaaaa tcatccgata ggaaaacttc gangaattgc 180  
tcnaantgct gggggtgaat gtgatgctna ngaanattgc tgtggctgca gcgtccaagc 240  
cagcagtggg gatcnaacag gagggagaca ctttctacat caaaacctcc accaccgtgc 300  
gcaccacaaa gattaacttc nnngttgggg aggantttga ggancaaact gtggatngga 360  
ngcctgtnaa aacctggtga aatgggagaa tganaataaa atggtctgtg ancanaaact 420  
cctgaaagga gaaggccccc anaactcctg gaccngaaaa actgaccnc cnatngggga 480  
actgatnctt gaaccctgaa cgggcgggat ganccttttt tnttgcncnc naanggggtc 540  
tttccntttc cccaaaaaaa 560

<210> 73  
<211> 379  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> 8, 17, 18, 21, 26, 29, 30, 32, 53, 56, 67, 71, 81, 102, 104,  
111, 112, 114, 119, 122, 124, 125, 134, 144, 146, 189, 190,  
214, 215, 219, 220, 235, 237, 246, 280, 288, 302, 310, 313,  
319, 322, 343, 353, 354  
<223> n = A,T,C or G

<400> 73  
ctggggancc ggcggtnngc nccatntcnn gncgcgaagg tggcaataaa aanccnctga 60  
aaccgcncaa naacatgcc naagatatgg acgaggaaga tngngctttc nngnacaanc 120  
gnanngagga acanaacaaa ctcnangagc tctcaagcta atgccgcggg gaagggggccc 180  
ttggccacnn gtggaattaa gaaatctggc aaanngtann tgttccttgt gcctnangag 240  
ataaangacc ctttattttca tctgtattta aacctctctn ttccctgnca taacttcttt 300  
tnccacgtan agntggaant anttgttgtc ttggactgtt gtncatttta gannaaactt 360  
ttgttcaaaa aaaaaataa 379

<210> 74  
<211> 437  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> 145, 355  
<223> n = A,T,C or G

&lt;400&gt; 74

```

actagttcag actgccacgc caaccccaga aaatacccca catgccagaa aagtgaagtc 60
ctaggtgttt ccactctatgt ttcaatctgt ccacttacca ggctctcgca taaaaacaaa 120
acaaaaaaac gctgccaggt ttanaagca gttctggctc caaaaccatc aggatcctgc 180
caccagggtt cttttgaaat agtaccacat gtaaaaggga atttggcttt cacttcatct 240
aatcactgaa ttgtcaggct ttgattgata attgtagaaa taagtagcct tctgttgg 300
gaataagtta taatcagtat tcatctcttt gttttttgtc actcttttct ctctnattgt 360
gtcatttgta ctgtttgaaa aatatttctt ctataaaatt aaactaacct gccttaaaaa 420
aaaaaaaaaa aaaaaaa 437

```

&lt;210&gt; 75

&lt;211&gt; 579

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; 440, 513, 539, 551

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 75

```

ctccgtcgcc gccaagatga tgtgcggggc gccctccgcc acgcagccgg ccaccgccga 60
gaccacagcac atcgccgacc aggtgaggct ccagcttgaa gagaaagaaa acaagaagtt 120
ccctgtgttt aaggccgtgt cattcaagag ccagggtgtc gcggggacaa actacttcat 180
caagggtcac gtcggcgacg aggacttctg acacctgcga gtgttccaat ctctccctca 240
tgaaaacaag cccttgacct tatctaacta ccagaccaac aaagccaagc atgatgagct 300
gacctatttc tgatcctgac tttggacaag gcccttcagc cagaagactg acaaagtcac 360
cctccgtcta ccagagcgtg cacttgtgat cctaaaataa gcttcatctc cgggctgtgc 420
ccttgggggtg gaagggggcan gatctgcact gcttttgcat ttctcttctt aaatttcatt 480
gtgttgattc tttccttcca atagggtgat ttnattactt tcagaatatt ttccaaatna 540
gatatatattt naaaatcctt aaaaaaaaaa aaaaaaaaaa 579

```

&lt;210&gt; 76

&lt;211&gt; 666

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; 411, 470, 476, 491, 506, 527, 560, 570, 632, 636, 643, 650, 654, 658

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 76

```

gtttatccta tctctccaac cagattgtca gctccttgag ggcaagagcc acagtatatt 60
tccctgtttc ttccacagtg cctaataata ctgtggaact aggttttaaat aatttttta 120
ttgatgttgt tatgggcagg atggcaacca gaccattgtc tcagagcagg tgctggctct 180
ttcctggcta ctccatgttg gctagcctct ggtaacctct tacttattat cttcaggaca 240
ctcactacag ggaccaggga tgatgcaaca tccttgtctt tttatgacag gatgtttgct 300
cagcttctcc aacaataaaa agcacgtggg aaaacacttg cggatattct ggactgtttt 360
taaaaaatat acagtttacc gaaaatcata ttatcttaca atgaaaagga ntttatagat 420
cagccagtga acaacctttt cccaccatac aaaaattcct tttcccgaa gaaaanggct 480
ttctcaataa ncctcacttt cttaanatct tacaagatag ccccganac ttatcgaaac 540
tcatttttagg caaatatgan ttttattgtg cgttacttgt ttcaaaattt ggtattgtga 600

```

```

atatcaatta ccacccccat ctcccatgaa anaaanggga aanggtgaan ttcntaancg 660
cttaaa 666

```

```

<210> 77
<211> 396
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 31, 54, 125, 128, 136, 163, 168, 198
<223> n = A,T,C or G

```

```

<400> 77
ctgcagcccg ggggatccac taatctacca nggttatttg gcagctaatt ctanatttgg 60
atcattgccc aaagttgcac ttgctgggtct ctggggattt ggccttggaagggtatcata 120
catanganta tgccanaata aattccattt ttttgaaaat canctccntg gggctgggtt 180
tgggtccacag cataacangc actgcctcct tacctgtgag gaatgcaaaa taaagcatgg 240
attaagtggag aagggagact ctgcgccttc agcttcctaa attctgtgtc tgtgactttc 300
gaagtttttt aaacctctga atttgtagac atttaaaatt tcaagtgtac tttaaaataa 360
aatacttcta atgggaacaa aaaaaaaaaa aaaaaa 396

```

```

<210> 78
<211> 793
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 309, 492, 563, 657, 660, 703, 708, 710, 711, 732, 740, 748,
758, 762, 765, 787
<223> n = A,T,C or G

```

```

<400> 78
gcacccctagc cgccgactca cacaaggcag gtgggtgagg aaatccagag ttgccatgga 60
gaaaattcca gtgtcagcat tcttgctcct tgtggccctc tcctacactc tggccagaga 120
taccacagtc aaacctggag ccaaaaagga cacaaggac tctcgacca aactgcccc 180
gacctctctc agagggtggg gtgaccaact catctggact cagacatatg aagaagctct 240
atataaatcc aagacaagca acaaaccctt gatgattatt catcacttgg atgagtggcc 300
acacagtcna gctttaaaga aagtgtttgc tgaaaaataaa gaaatccaga aattggcaga 360
gcagtttgtc ctctcaatc tggtttatga aacaactgac aaacaccttt ctctgatgg 420
ccagtatgtc ccaggattat gtttgttgac ccatctctga cagttgaagc cgatatcctg 480
ggaagatatt cnaaccgtct ctatgcttac aaactgcaga tacgctctgt tgcttgacac 540
atgaaaaagc tctcaagttg ctnaaaatga attgtaagaa aaaaaatctc cagccttctg 600
tctgtcggct tgaaaaattga aaccagaaaa atgtgaaaaa tggctattgt ggaacanatn 660
gacacctgat taggttttgg ttatgttcac cactattttt aanaaaanan nttttaaaat 720
ttggttcaat tntctttttn aaacaatntg tttctacntt gnganctgat ttctaaaaaa 780
aataatnttt ggc 793

```

```

<210> 79
<211> 456
<212> DNA
<213> Homo sapiens

```

<220>  
 <221> misc\_feature  
 <222> 89, 195, 255, 263, 266, 286, 353, 384, 423, 425, 436, 441  
 <223> n = A,T,C or G

<400> 79  
 actagtatgg ggtgggaggc cccacccttc tcccctaggc gctgttcttg ctccaaaggg 60  
 ctccgtggag agggactggc agagctgang ccacctgggg ctggggatcc cactcttctt 120  
 gcagctgttg agcgcaccta accactggtc atgccccac ccctgctctc cgcacccgct 180  
 tcctcccgac cccangacca ggctacttct cccctcctct tgccctccctc ctgcccctgc 240  
 tgcctctgat cgtangaatt gangantgtc ccgccttgtg gctganaatg gacagtggca 300  
 ggggctggaa atgggtgtgt gtgtgtgtgt gtgtgtgtgt gtgtgtgtgt gcnccccccc 360  
 tgcaagaccg agattgaggg aaancatgtc tgctgggtgt gacctgttt cctctccata 420  
 aantncccct gtgacnctca naaaaaaaaa aaaaaa 456

<210> 80  
 <211> 284  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 283  
 <223> n = A,T,C or G

<400> 80  
 ctttgtacct ctagaaaaga taggtattgt gtcataaaac ttgagtttaa attttatata 60  
 taaaactaaa agtaatgctc acttttagcaa cacatactaa aattggaacc atactgagaa 120  
 gaatagcatg acctccgtgc aaacaggaca agcaaatttg tgatgtgttg attaaaaaga 180  
 aataaataaa tgtgtatatg tgtaacttgt atgtttatgt ggaatacaga ttgggaaata 240  
 aaatgtattt cttactgtga aaaaaaaaaa aaaaaaaaaa aana 284

<210> 81  
 <211> 671  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 388, 505, 600, 603, 615, 642, 644, 660  
 <223> n = A,T,C or G

<400> 81  
 gccaccaaca ttccaagcta ccctgggtac ctttgtgcag tagaagctag tgagcatgtg 60  
 agcaagcggg gtgcacacgg agactcatcg ttataattta ctatctgcca agagtagaaa 120  
 gaaaggctgg ggatatttgg gttggcttgg ttttgatttt ttgcttggtt gtttgttttg 180  
 tactaaaaca gtattatctt ttgaatatcg tagggacata agtatataca tgttatccaa 240  
 tcaagatggc tagaatgggtg cctttctgag tgtctaaaac ttgacacccc tggtaaatct 300  
 ttcaacacac ttccactgcc tgcgtaatga agttttgatt catttttaac cactggaatt 360  
 tttcaatgcc gtcattttca gttagatnat tttgcacttt gagattaaaa tgccatgtct 420  
 atttgattag tcttattttt ttattttttac aggcttatca gtctcactgt tggctgtcat 480  
 tgtgacaaaag tcaataaaac cccnaggac aacacacagt atgggatcac atattgtttg 540  
 acattaagct ttggccaaaa aatgttgcac gtgtttttacc tgcacttgct aaatcaatan 600  
 canaaaggct ggctnataat gttgggtgggtg aaataattaa tnantaacca aaaaaaaaaa 660

aaaaaaaaa a

671

<210> 82  
<211> 217  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> 35  
<223> n = A,T,C or G

<400> 82  
ctgcagatgt ttcttgaatg ctttgtcaaa ttaanaaagt taaagtgcaa taatgtttga 60  
agacaataag tgggtggtgta tcttgtttct aataagataa acctttttgt ctttgcttta 120  
tcttattagg gagttgtatg tcagtgtata aaacatactg tgtggtataa caggcttaat 180  
aaattcttta aaaggaaaaa aaaaaaaaaa aaaaaaa 217

<210> 83  
<211> 460  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> 104, 118, 172, 401, 422, 423, 444, 449  
<223> n = A,T,C or G

<400> 83  
cgcgagtggg agcaccagga tctcggggctc ggaacgagac tgcacggatt gttttaagaa 60  
aatggcagac aaaccagaca tgggggaaat cgccagcttc gatnaggcca agctgaanaa 120  
aacggagacg caggagaaga acaccctgcc gaccaaagag accattgagc angagaagcg 180  
gagtgaattt tcctaagatc ctggaggatt tcctaccccc gtctctctcg agaccccagt 240  
cgtgatgtgg aggaagagcc acctgcaaga tggacacgag ccacaagctg cactgtgaac 300  
ctgggcactc cgcgccgatg ccaccggcct gtgggtctct gaagggaccc cccccaatcg 360  
gactgccaaa ttctccggtt tgccccggga tattatacaa nattatttgt atgaataatg 420  
annataaaac acacctcgtg gcancaaana aaaaaaaaaa 460

<210> 84  
<211> 323  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> 70, 138, 178, 197, 228, 242, 244, 287, 311  
<223> n = A,T,C or G

<400> 84  
tgggtgatct tggctctgtg gagctgctgg gacgggatct aaaagactat tctggaagct 60  
gtggtccaan gcattttgct ggcttaacgg gtcccgaac aaaggacacc agctctctaa 120  
aattgaagtt tacccganat aacaatcttt tgggcagaga tgcttatatt aacaaacncc 180  
gtccctgcgc aacaacnaac aatctctggg aaataccggc catgaacntg ctgtctcaat 240  
cnancatctc tctagctgac cgatcatatc gtcccagatt actacanatc ataataattg 300



atttcctgta naaaaaaaaa aaa

323

<210> 85  
<211> 771  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> 63, 426, 471, 497, 521, 554, 583, 586, 606, 609, 615, 652,  
686, 691, 694, 695, 706, 713, 730, 732, 743, 751  
<223> n = A,T,C or G

<400> 85  
aaactgggta ctcaacactg agcagatctg ttctttgagc taaaaacccat gtgctgtacc 60  
aanagtttgc tcctggctgc tttgatgtca gtgctgctac tccacctctg cggcgaaatca 120  
gaagcaagca actttgactg ctgtcttggg tacacagacc gtattcttca tcctaaattt 180  
attgtgggct tcacacggca gctggccaat gaaggctgtg acatcaatgc tatcatcttt 240  
cacacaaaga aaaagttgtc tgtgtgcgca aatccaaaac agacttgggt gaaatatatt 300  
gtgctgtctc tcagtaaaaa agtcaagaac atgtaaaaac tgtggctttt ctggaatgga 360  
attggacata gcccaagaac agaaagaact tgctgggggt ggagggttca ctgacacatc 420  
atgganggtt tagtgcttat cttattttgtg cctcctggac ttgtccaatt natgaagtta 480  
atcatattgc atcatanttt gctttgttta acatcacatt naaatttaaac tgtattttat 540  
gttattttata gctntaggtt ttctgtgttt aactttttat acnaantttc ctaaactatt 600  
ttggtntant gcaanttaaa aattatattt ggggggggaa taaatattgg antttctgca 660  
gccacaagct ttttttaaaa aaccantaca nccnngttaa atggtnngtc ccnaatgggt 720  
tttgcttttn antagaaaat ttnttiagaac natttgaaaa aaaaaaaaaa a 771

<210> 86  
<211> 628  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> 162, 249, 266, 348, 407, 427, 488, 518, 545, 566, 569, 597,  
598, 611, 617, 621, 624  
<223> n = A,T,C or G

<400> 86  
actagtttgc tttacatttt tgaaaagtat tatttttgtc caagtgttta tcaactaaac 60  
cttgtgttag gtaagaatgg aatttattaa gtgaatcagt gtgacccttc ttgtcataag 120  
attatcttaa agctgaagcc aaaatatgct tcaaaagaaa angactttat tgttcattgt 180  
agttcataca ttcaaagcat ctgaactgta gtttctatag caagccaatt acatccataa 240  
gtggagaang aaatagatta atgtcnaagt atgattgggt gagggagcaa gggtgaagat 300  
aatctggggt tgaaattttc tagttttcat tctgtacatt tttagttnga catcagattt 360  
gaaatattaa tgtttacctt tcaatgtgtg gtatcagctg gactcantaa cacccttttc 420  
ttccctnngg gatggggaat ggattattgg aaaatggaaa gaaaaaagta cttaaagcct 480  
tcctttcnca gtttctggct cctaccctac tgatttancc agaataagaa aacattttat 540  
catcntctgc tttattccca ttaatnaant tttgatgaat aaatctgctt ttatgcnnac 600  
ccaaggaatt nagtggnttc ntcnttgt 628

<210> 87  
<211> 518

<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> 384, 421, 486  
<223> n = A,T,C or G

<400> 87  
 ttttttatttt ttttttagaga gtagttcagc tttttatttat aaattttattg cctgtttttat 60  
 tataacaaca ttatactggt tatggtttaa tacatatggt tcaaaatgta taatacatca 120  
 agtagtacag ttttaaaatt ttatgcttaa aacaagtttt gtgtaaaaaa tgcagatata 180  
 ttttacatgg caaatcaatt ttttaagtcac cctaaaaatt gatttttttt tgaaatttaa 240  
 aaacacatttt aattttcaatt tctctcttat ataacccttta ttactatagc atgggtttcca 300  
 ctacagttta acaatgcagc aaaattccca tttcacggta aattggggtt taagcggcaa 360  
 gggttaaaatg ctttgaggat cctnaatacc ctttgaactt caaatgaagg ttatgggtgt 420  
 naatttaacc ctcatgccat aagcagaagc acaagtttag ctgcattttg ctctaaactg 480  
 taaaancgag ccccccggtg aaaaagcaaa aggggaccc 518

<210> 88  
<211> 1844  
<212> DNA  
<213> Homo sapiens

<400> 88  
 gagacagtga atcctagtagt caaaggatttt ttggcctcag aaaaagttgt tgattattttt 60  
 tattttatttt tattttttcga gactccgtct caaaaaaaaaa aaaaaaaaaa agaatacaca 120  
 ggtatttgct aaagcatttt gagctgcttg gaaaaaggga agtagttgca gtagagtttc 180  
 ttccatcttc ttggtgctgg gaagccatat atgtgtcttt tactcaagct aaggggtata 240  
 agcttatgtg ttgaattttgc tacatctata ttccacatat tctcacaata agagaatttt 300  
 gaaatagaaa tatcatagaa catttaagaa agtttagtat aaataatatt ttgtgtgttt 360  
 taatcccttt gaagggatct atccaaagaa aatattttac actgagctcc ttccacacg 420  
 tctcagtaac agatcctgtg ttagtctttg aaaatagctc atttttttaa tgtcagtgag 480  
 tagatgtagc atacatatga tgtataatga cgtgtattat gttaacaatg tctgcagatt 540  
 ttgtaggaat acaaaacatg gcctttttta taagcaaaac gggccaatga ctagaataac 600  
 acatagggca atctgtgaat atgtattata agcagcattc cagaaaagta gttggtgaaa 660  
 taattttcaa gtcaaaaagg gatattgaaa gggaattatg agtaacctct attttttaag 720  
 ccttgctttt aaattaaacg ctacagccat ttaagccttg aggataataa agcttgagag 780  
 taataatggt aggttagcaa aggttttagat gtatcacttc atgcatgcta ccatgatagt 840  
 aatgcagctc ttcgagtcac ttctggtcac tcaagatatt cacccttttg cccatagaaa 900  
 gcaccctacc tcacctgctt actgacattg tcttagctga tcacaagatc attatcagcc 960  
 tccattattc cttactgtat ataaaaataca gagttttata ttttcctttc ttctgttttc 1020  
 accatattca aaacctaaat ttgtttttgc agatggaatg caaagtaatc aagtgttcgt 1080  
 gctttcacct agaagggtgt ggtcctgaag gaaagaggtc cctaaatatc cccacacctg 1140  
 ggtgctctc cttccctggt accctgacta ccagaagtca ggtgctagag cagctggaga 1200  
 agtgcagcag cctgtgcttc cacagatggg ggtgctgctg caacaaggct ttcaatgtgc 1260  
 ccattcttag gggagaagct agatcctgtg cagcagcctg gtaagtcctg aggaggttcc 1320  
 attgctcttc ctgctgctgt ccttttgcttc tcaacggggc tcgctctaca gtctagagca 1380  
 catgcagcta acttgtgcct ctgcttatgc atgaggggta aattaacaac cataaccttc 1440  
 atttgaagtt caaagggtgta ttcaggatcc tcaaagcatt ttaaccttgc cgcttaaaac 1500  
 ccaatttacc gtgaaatggg aattttgctg cattgtttaa ctgtagtgga aaccatgcta 1560  
 tagtaataaa ggttatataa gagagaaatt gaaattaaat gtgtttttta atttcaaaaa 1620  
 aaaatcaatc tttaggatga cttaaaaaatt gatttgccat gtaaaatgta tctgcatttt 1680  
 ttacacaaaa cttgttttaa gcataaaaatt ttaaaactgt actacttgat gtattatata 1740

```

ttttgaacca tatgtattaa accataaaca gtataatggt gttataataa aacaggcaat 1800
aaattttataa ataaaaagctg aaaaaaaaaa aaaaaaaaaa aaaa 1844

```

```

<210> 89
<211> 523
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 288, 352, 369, 398, 475, 511, 513
<223> n = A,T,C or G

```

```

<400> 89
tttttttttt ttttttttagt caatccacat ttattgatca cttattatgt accaggcact 60
gggataaaaga tgactgttag tcaactcacag taagggaagaa aactagcaaa taagacgatt 120
acaatatgat gtagaaaatg ctaagccaga gatatagaaa ggtcctattg ggtccttctg 180
tcacctgtgc ttccacacat cctacccttc acaggccttc cctccagctt cctgcccccg 240
ctccccactg cagatcccct gggattttgc ctgagagctaa acgagganat gggccccctg 300
gccctggcat gacttgaacc caaccacaga ctgggaaagg gagcctttcg anagtggatc 360
actttgatna gaaaacacat agggaattga agagaaantc cccaaatggc caccctgtgct 420
gggtgtcaag aaaagtttgc agaattggata aatgaaggat caagggaatt aatanatgaa 480
taattgaatg gtggctcaat aagaatgact ncnttgaatg acc 523

```

```

<210> 90
<211> 604
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 563
<223> n = A,T,C or G

```

```

<400> 90
ccagtgtggt ggaatgcaaa gattaccccg gaagctttcg agaagctggg attccctgca 60
gcaaaggaaa tagccaatat gtgtcgtttc tatgaaatga agccagaccg agatgtcaat 120
ctcaccacc aactaaatcc caaagtcaaa agcttcagcc agtttatctc agagaaccag 180
gggagccttc aaggggcatgt agaaaatcag ctgttcagat aggcctctgc accacacagc 240
ctctttcctc tctgacacct ttctcttcta cggcacaaca ttcattgtttg acagaacatg 300
ctggaatgca attgtttgca acaccgaagg atttcctgcg gtcgcctctt cagtaggaag 360
cactgcattg gtgataggac acggtaatat gattcacatt taacttgcta gttagtata 420
aggggtggta cacctgtttg gtaaaatgag aagcctcgga aacttgggag cttctctcct 480
accactaatg gggagggcag attattactg ggatttctcc tgggggtgaat taatttcaag 540
ccctaattgc tgaaattccc ctnggcaggc tccagttttc tcaactgcat tgcaaaattc 600
cccc 604

```

```

<210> 91
<211> 858
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature

```

<222> 570, 591, 655, 664, 667, 683, 711, 759, 760, 765, 777, 787,  
792, 794, 801, 804, 809, 817, 820

<223> n = A,T,C or G

<400> 91

```

tttttttttt ttttttttta tgattattat ttttttttatt gatctttaca tcctcagtgt 60
tggcagagtt tctgatgctt aataaacatt tgttctgata agataagtgg aaaaaattgt 120
catttcctta ttcaagccat gctttttctgt gatattctga tcctagttga acatacagaa 180
ataaatgtct aaaacagcac ctcgattctc gtctataaca ggactaagtt cactgtgatc 240
ttaaataagc ttggctaaaa tgggacatga gtggaggtag tcacacttca gcgaagaaag 300
agaatctcct gtataatctc accaggagat tcaacgaatt ccaccacact ggactagtgg 360
atcccccggt ctgcaggaat tcgatatcaa gcttatcgat accgtcgacc tcgagggggg 420
gcccgttacc caattcgccc tatagttagt cgtattacgc gcgctcactg gccgtcgttt 480
tacaacgtcg tgactgggaa aaccctggcg ttacccaact taatcgccct gcagcacatc 540
cccctttcgc cagctggcgt aatagcgaan agcccgcacc gatcgccctt ncaacagttg 600
cgcagcctga atggcgaaat ggacgcgccc tgtagcggcg cattaaagcg cggcnggggtg 660
tggnggntcc cccacgtgac cgntacactt ggcagcgccct tacgcccgtc ntctcgctttc 720
ttcccttcct ttctcgacc gttcgccggg tttccccggn agctnttaat cgggggnctc 780
cctttanggg tncnaattaa nggnttacng gaccttngan cccaaaaact ttgattaggg 840
ggaaggtccc cgaagggg                                     858

```

<210> 92

<211> 585

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 317, 319, 320, 321, 325, 327, 328, 330, 331, 332, 460, 462,  
483, 485, 487, 523, 538, 566, 584

<223> n = A,T,C or G

<400> 92

```

gttgaatctc ctggtgagat tatacaggag attctctttc ttcgctgaag tgtgactacc 60
tccactcatg tcccatttta gccaaagctta tttaagatca cagtgaactt agtcctgtta 120
tagacgagaa tcgaggtgct gtttttagaca tttatttctg tatgttcaac taggatcaga 180
atatcacaga aaagcatggc ttgaataagg aaatgacaat tttttccact tatctgatca 240
gaacaaatgt ttattaagca tcagaaactc tgccaacact gaggatgtaa agatcaataa 300
aaaaataaat aatcatnann naaanannan nngaaggggcg gccgccaccg cgggtggagct 360
ccagcttttg ttcccttttag tgagggttaa ttgcgcgctt ggcgttaatc atgggtcatag 420
ctgtttcctg tgtgaaattg ttatccggct cacaattccn cncaacatac gagccgggaa 480
gcntnangtg taaaagcctg ggggtgccta attgagtgag ctnactcaca ttaattgngt 540
tgcgctccac ttgcccgtt ttccantccg ggaaacctgt tcgnc                                     585

```

<210> 93

<211> 567

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 82, 158, 230, 232, 253, 266, 267, 268, 269, 270, 271, 272,  
273, 274, 275, 276, 277, 278, 279, 280, 281, 282, 283, 284,  
285, 286, 287, 295, 303, 307, 314, 349, 352, 354, 356, 366,

369, 379, 382, 386, 393, 404, 427, 428, 446, 450, 452

<223> n = A,T,C or G

<221> misc\_feature

<222> 453, 454, 459, 462, 480, 481, 483, 488, 493, 501, 509, 511, 512, 518, 520, 525, 526, 532, 541, 557

<223> n = A,T,C or G

<400> 93

```
cggcagtgtt gctgtctgcg tgtccacctt ggaatctggc tgaactggct gggaggacca 60
agactgcggc tggggtgggc anggaaggga accgggggct gctgtgaagg atcttggaac 120
ttccctgtac ccaccttccc cttgcttcat gtttgtanag gaaccttgtg ccggccaagc 180
ccagtttcct tgtgtgatac actaatgtat ttgctttttt tgggaaatan anaaaaatca 240
attaaattgc tantgtttct ttgaannnnn nnnnnnnnnn nnnnnnnggg ggggncgccc 300
ccncgngnga aacnccccct ttgtttccct ttaattgaaa ggttaattng cncncntggc 360
gttaanccnt gggccaaanc tngttncccg tgntgaaatt gttnatcccc tcccaaattc 420
cccccnccc ttccaaaccc ggaaanccctn annntgttna anccccgggg gttgcctaen 480
ngnaattnaa ccnaaccccc ntttaaattng nntttgcncn ccacnngccc cncctttccca 540
nttcggggaa aacctnttcc gtgccca 567
```

<210> 94

<211> 620

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 169, 171, 222, 472, 528, 559, 599

<223> n = A,T,C or G

<400> 94

```
actagtcaaa aatgctaaaa taatttgagg gaaaatattt tttaaagtagt gttatagttt 60
catgtttatc ttttattatg ttttgtgaag ttgtgtcttt tcactaatta cctatactat 120
gccaatattt ctttatatct atccataaca tttatactac atttgtaana naatatgcac 180
gtgaaactta acactttata aggtaaaaat gaggtttcca anatttaata atctgatcaa 240
gttcttgtaa tttccaaata gaatggactt ggtctgttaa gggctaagga gaagaggaag 300
ataagggtta aagttgttaa tgaccaaaca ttctaaaaga aatgcaaaaa aaaagtttat 360
tttcaagcct tcgaactatt taaggaaaagc aaaatcattt cctaaatgca tatcatttgt 420
gagaatttct cattaatatc ctgaatcatt catttcacta aggcctcatgt tnactccgat 480
atgtctctaa gaaagtacta tttcatggtc caaacctggg tgccatantt gggtaaaggc 540
tttcccttaa gtgtgaaant atttaaaatg aaattttcct ctttttaaaa attctttana 600
aggggttaagg gtgttgaggga 620
```

<210> 95

<211> 470

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 61, 67, 79, 89, 106, 213, 271, 281, 330, 354, 387, 432, 448

<223> n = A,T,C or G

<400> 95

```

ctcgaccttc tctgcacagc ggatgaaccc tgagcagctg aagaccagaa aagccactat 60
nactttntgc ttaattcang agcttacang attcttcaaa gagtgngtcc agcatccttt 120
gaaacatgag ttcttaccag cagaagcaga cctttacccc accacctcag cttcaacagc 180
agcaggtgaa acaacccatc cagcctccac ctnaggaaat atttgttccc acaaccaagg 240
agccatgcc a ctcaaagggtt ccacaacctg naaacacaaa nattccagag ccaggctgta 300
ccaaggtccc tgagccaggg ctgtaccaan gtccctgagc caggttgtac caangtccct 360
gagccaggat gtaccaagggt ccctgancca ggttggtccaa ggtccctgag ccaggctaca 420
ccaagggcct gngccaggca gcatcaangt ccctgaccaa ggcttatcaa 470

```

<210> 96

<211> 660

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 299, 311, 360, 426, 538, 540, 542, 553, 563, 565, 592, 603, 604, 618, 633, 647, 649, 651, 653

<223> n = A,T,C or G

<400> 96

```

tttttttttt tttttttttt ggaattaaaa gcaatttaat gagggcagag caggaaacat 60
gcatttcttt tcattcgaat cttcagatga accctgagca gccgaagacc agaaaagcca 120
tgaagacttt ctgcttaatt caggggctta caggattctt cagagtgtgt gtgaacaaaa 180
gctttatagt acgtattttt aggatacaaa taagagagag actatggctt ggggtgagaa 240
tgtactgatt acaagggtcta cagacaatta agacacagaa acagatggga agaggggtgnc 300
cagcatctgg nggttggctt ctcaagggct tgtctgtgca ccaaattact tctgcttgnn 360
cttctgctga gctgggcctg gagtgaacct tgaaggacat ggctctggta cctttgtgta 420
gcctgncaca ggaactttgg tgtatccttg ctcaggaaact ttgatggcac ctggctcagg 480
aaacttgatg aagccttggt caagggacct tgatgcttgc tggctcaggg accttggnn 540
ancctgggct canggacctt tgnncnaacc ttggcttcaa gggacccttg gnacatcctg 600
gcnnagggac ccttgggncc aaccctgggc ttnagggacc ctttggnntnc nanccttggc 660

```

<210> 97

<211> 441

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 12, 308

<223> n = A,T,C or G

<400> 97

```

gggaccatac anagtattcc tctcttcaca ccaggaccag ccactgttgc agcatgagtt 60
cccagcagca gaagcagccc tgcatccac cccctcagct tcagcagcag cagggtgaaac 120
agccttgcca gctccacct caggaacct gcaccccaa aaccaaggag ccctgccacc 180
ccaaggtgcc tgagccctgc cccccaaag tgccctgagcc ctgccagccc aaggttccag 240
agccatgcc ccccaagggtg cctgagccct gcccttcaat agtcaactca gcaccagccc 300
agcagaanac caagcagaag taatgtggct cacagccatg cccttgagga gccggccacc 360
agatgctgaa tcccctatcc cattctgtgt atgagtccca tttgccttgc aattagcatt 420
ctgtctcccc caaaaaaaaa a

```

<210> 98  
 <211> 600  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 295, 349, 489, 496, 583  
 <223> n = A,T,C or G

<400> 98  
 gtatttcctct cttcacacca ggaccagcca ctgttgccagc atgagttccc agcagcagaa 60  
 gcagccctgc atcccccccc ctcagcttca gcagcagcag gtgaaacagc cttgccagcc 120  
 tccacctcag gaacctatgca tccccaaaac caaggagccc tgccacccca aggtgcctga 180  
 gccctgccac cccaaagtgc ctgagccctg ccagcccagc gttccagagc catgccaccc 240  
 caaggtgcct gagccctgcc cttcaatagt cactccagca ccagcccagc agaanaccaa 300  
 gcagaagtaa tgtgggtccac agccatgccc ttgaggagcc ggccaccana tgctgaatcc 360  
 cctatcccat tctgtgtatg agtcccattt gccttgcaat tagcattctg tctcccccaa 420  
 aaaagaatgt gctatgaagc tttctttcct acacactctg agtctctgaa tgaagctgaa 480  
 ggtcttaant acaganctag ttttcagctg ctcagaattc tctgaagaaa agattttaaga 540  
 tgaaaggcaa atgattcagc tccttattac cccattaaat tcncctttcaa ttccaaaaaa 600

<210> 99  
 <211> 667  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 345, 562, 635  
 <223> n = A,T,C or G

<400> 99  
 actagtgact gagttcctgg caaagaaatt tgacctggac cagttgataa ctcatgtttt 60  
 accattttaa aaaatcagtg aaggatttga gctgctcaat tcaggacaaa gcattcgaac 120  
 ggtcctgacg ttttgagatc caaagtggca ggaggctctg gttgtcatgg tgaactggag 180  
 tttctcttgt gagagttccc tcatctgaaa tcatgtatct gtctcacaaa tacaagcata 240  
 agtagaagat ttgttgaaga catagaaccc ttataaagaa ttattaacct ttataaacat 300  
 ttaaagtctt gtgagcacct gggaattagt ataataacaa tgttnatatt tttgatttac 360  
 attttgtaag gctataattg tatcttttaa gaaaacatac cttggatttc tatgttgaaa 420  
 tggagatttt taagagtttt aaccagctgc tgcagatata ttactcaaaa cagatatagc 480  
 gtataaagat atagtaaagc catctcctag agtaatatc acttaacaca ttggaaacta 540  
 ttatttttta gatttgaata tnaatgttat tttttaaaca cttgttatga gttacttggg 600  
 attacatttt gaaatcagtt cattccatga tgcanattac tgggattaga ttaagaaaaga 660  
 cggaataa 667

<210> 100  
 <211> 583  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature

<222> 404, 506, 514, 527, 528, 538, 548, 556, 568, 569

<223> n = A,T,C or G

<400> 100

```
gttttgtttg taagatgata acagtcattgt tacactgata taaaggacat atatataacc 60
ctttaaaaaa aaaatcactg cctcattctt atttcaagat gaatttctat acagactaga 120
tgtttttctg aagatcaatt agacattttg aaaatgattt aaagtgtttt ccttaatgtt 180
ctctgaaaac aagtttcttt tgtagtttta accaaaaaag tgcccttttt gtcactggat 240
tctcctagca ttcattgattt ttttttcata caatgaaatt aaaatttgta aaatcatgga 300
ctggctttct gggtggattt caggtaagat gtgtttaagg ccagagcttt tctcagtatt 360
tgattttttt ccccaatatt tgatttttta aaaatataca catnggtgct gcattttatat 420
ctgctggttt aaaattctgt catatttcac ttctagcctt ttagttatgg caaatcatat 480
tttactttta cttaaagcat ttggttattt ggantatctg gttctannct aaaaaanta 540
attctatnaa ttgaantttt ggtactcnnn catatttgga tcc 583
```

<210> 101

<211> 592

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 218, 497, 502, 533, 544, 546, 548, 550, 555

<223> n = A,T,C or G

<400> 101

```
gtggagacgt acaaagagca gccgctcaag acacctggga agaaaaagaa aggcaagccc 60
gggaaacgca aggagcagga aaagaaaaaa cggcgaactc gctctgcctg gttagactct 120
ggagtgcact ggagtgggct agaaggggac cacctgtctg acacctccac aacgtcgctg 180
gagctcgatt caccgaggca ttgaaatttt cagcaganac cttccaagga catattgcag 240
gattctgtaa tagtgaacat atggaaagta ttagaaatat ttattgtctg taaatactgt 300
aaatgcattg gaataaaact gtctccccc ttgctctatg aaactgcaca ttgggtcattg 360
tgaatatatt tttttttgcc aaggctaata caattattat tatcacattt accataattt 420
attttgtcca ttgatgtatt tattttgtaa atgtatcttg gtgctgctga atttctatat 480
tttttgtaca taatgcnitt anatatacct atcaagtttg ttgataaatg acncaatgaa 540
gtgncncnan ttgngnggtt aatttaatga atgcctaatt ttattatccc aa 592
```

<210> 102

<211> 587

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 91, 131, 256, 263, 332, 392, 400, 403, 461, 496, 497, 499, 510, 511, 518, 519, 539, 554, 560, 576

<223> n = A,T,C or G

<400> 102

```
cgtcctaagc acttagacta catcagggaa gaacacagac cacatccctg tcctcatgct 60
gcttatgttt tctggaagaa agtggagacc nagtccttgg ctttagggct ccccggtctg 120
gggctgtgca ntccggctcag ggcgggaagg gaaatgcacc gctgcatgtg aacttacagc 180
ccaggcggat gcccttccc ttagcactac ctggcctcct gcacccctc gcctcatgtt 240
cctccacact tcaanaaatg aanaacccca tgggcccagc cccttgccct ggggaaccaa 300
```



```

ggcagccttc caaaaactcag gggctgaagc anactattag ggcaggggct gactttgggt 360
gacactgccc attccctctc agggcagctc angtcacccn ggncctctga acccagcctg 420
ttcctttgaa aaaggggcaaa actgaaaagg gcttttccta naaaaagaaa aaccagggaa 480
ctttgccagg gcttcnntnt taccaaaacn ncttctcnng gatttttaat tccccatng 540
gcctccactt accnngggcn atgccccaaa attaanaatt tcccatc 587

```

```

<210> 103
<211> 496
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 2, 17, 66, 74, 82, 119, 164, 166, 172, 200, 203, 228, 232,
271, 273, 415, 423, 445, 446, 473
<223> n = A,T,C or G

```

```

<400> 103
anaggactgg ccctacntgc tctctctcgt cctacctatc aatgccaac atggcagaac 60
ctgcanccct tggncactgc anatggaaac ctctcagtggt cttgacatca ccctaccnt 120
gcggtgggtc tccaccacaa ccactttgac tctgtggtcc ctgnanggtg gnttctcctg 180
actggcagga tggaccttan ccnecatatc cctctgttcc ctctgctnag anaaagaatt 240
cccttaacat gatataatcc acccatgcaa ntngctactg gccagctac catttaccat 300
ttgcctacag aatttcattc agtctacact ttggcattct ctctggcgat agagtgtggc 360
tgggctgacc gcaaaaagggtg ccttacacac tggccccac cctcaaccgt tgacncatca 420
gangcttgcc tctccttct gattnncccc catgttggat atcaggggtg tcnagggatt 480
ggaaaaagaaa caaaac 496

```

```

<210> 104
<211> 575
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 18, 19, 45, 68, 77, 132, 155, 174, 219, 226, 238, 259, 263,
271, 273, 306, 323, 339, 363, 368, 370, 378, 381, 382, 436,
440, 449, 450, 456, 481, 485, 496, 503, 510, 512, 515, 528,
542, 552
<223> n = A,T,C or G

```

```

<400> 104
gcacctgctc tcaatccnnc tctcaccatg atcctccgcc tgcanaaact cctctgcca 60
ctatggangt ggtttcnngg gtggctcttg ccaactggga agaagccgtg gtgtctctac 120
ctgttcaact cngtttgtgt ctggggggtc aactnngggc tatggaagcg gctnaactgt 180
tgttttggtg gaagggtctg taattggctt tgggaagtng cttatngaag ttggcctnng 240
gaagttgcta ttgaaagtng ccntggaaagt ngntttggtg gggggttttg ctggtggcct 300
ttgttnaatt tgggtgcttt gtnaatggcg gccccctcnc ctgggcaatg aaaaaaatca 360
ccnatgcngn aaacctcnac nnaacagcct gggcttccct cacctcgaaa aaagtgtgctc 420
cccccccaaa aaaggncaan cccctcaann tggangttg aaaaaatcct cgaatgggga 480
nccnnaaaac aaaaancccc cnttttccn gnaanggggg aaataccncc cccccactta 540
cnaaaaccct tntaaaaaac cccccgggaa aaaaa 575

```

```

<210> 105

```

<211> 619  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 260, 527, 560, 564, 566, 585, 599  
 <223> n = A,T,C or G

<400> 105  
 cactagtagg atagaaacac tgtgtcccgag gagtaaggag agaagctact attgattaga 60  
 gcctaaccga ggtaactgc aagaagaggc gggatacttt cagctttcca tgtaactgta 120  
 tgcataaagc caatgtagtc cagtttctaa gatcatgttc caagctaact gaatcccact 180  
 tcaatacaca ctcatgaact cctgatggaa caataacagg cccaagcctg tggatgatg 240  
 tgcacacttg ctgactcan aaaaaatact actctcataa atgggtggga gtattttggg 300  
 gacaacctac tttgcttggc tgagtgaagg aatgatattc atatattcat ttattccatg 360  
 gacatttagt tagtgctttt tatataccag gcatgatgct gagtgacact cttgtgtata 420  
 tttccaaatt tttgtacagt cgctgcacat atttgaaatc atatattaag acttccaaaa 480  
 aatgaagtcc ctggtttttc atggcaactt gatcagtaaa ggattcncct ctgttttggt 540  
 cttaaaacat ctactatatn gttanatga aattcctttt ccccnctcc cgaaaaaana 600  
 aagtgtggg gaaaaaaaaa 619

<210> 106  
 <211> 506  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 8, 21, 31, 32, 58, 75, 89, 96, 99, 103, 122, 126, 147, 150,  
 158, 195, 210, 212, 219, 226, 246, 248, 249, 255, 258, 261,  
 263, 265, 275, 304, 317, 321, 331, 337, 340, 358, 371, 377,  
 380, 396, 450, 491  
 <223> n = A,T,C or G

<400> 106  
 cattggtinct ttcatttgct ntggaagtgt nnatctctaa cagtggacaa agttcccngt 60  
 gccttaaaact ctgtnacact tttgggaant gaaaanttng tantatgata ggttattctg 120  
 angtanagat gttctggata ccattanatn tgcccccnct gtcagaggct catatttgtt 180  
 tatgtaaatg gtatntcatt cgctactatn antcaattng aaatanggtc tttgggttat 240  
 gaatantnng cagencanct nananctgt ctgtngtatt cattgtggtc atagcacctc 300  
 acancattgt aacctcnatc nagtgagaca nactagnaan ttcctagtga tggctcanga 360  
 ttccaaatgg nctcatntcn aatgttttaa agttanttaa gtgtaagaaa tacagactgg 420  
 atgttccacc aactagtacc tgtaatgacn ggctgtccc aacacatctc ccttttccat 480  
 gactgtggtgta nccgcgcatcg gaaaaa 506

<210> 107  
 <211> 452  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 289, 317, 378

<223> n = A,T,C or G

<400> 107

```
gttgagtctg tactaaacag taagatatct caatgaacca taaattcaac tttgtaaaaa 60
tcttttgaag catagataat attgttttgt aaatgtttct tttgttttgt aaatgtttct 120
tttaaagacc ctcctattct ataaaaactct gcatgtagag gcttggttac ctttctctct 180
ctaaggttta caataggagt ggtgatttga aaaatataaa attatgagat tgggttttcct 240
gtggcataaa ttgcatcact gtatcatttt ctttttttaac cggtaagant ttcagtttgt 300
tggaaaagtaa ctgtganaac ccagtttccc gtccatctcc cttagggact acccatagaa 360
catgaaaagg tccccacnga agcaagaaga taagtctttc atggctgctg gttgcttaaa 420
ccacttttaa accaaaaaat tccccttggg aa 452
```

<210> 108

<211> 502

<212> DNA

<213> Homo sapiens

<220>

<221> misc feature

<222> 22, 31, 126, 168, 183, 205, 219, 231, 236, 259, 283, 295,  
296, 298, 301, 340, 354, 378, 383, 409, 433, 446, 455, 466,  
488

<223> n = A,T,C or G

<400> 108

```
atcttcttcc cttaattagt tnttatattat ntattaaatt ttattgcatg tcctggcaaa 60
caaaaaagaga ttgtagattg gcttctggct ccccaaaagc ccataacaga aagtaccaca 120
agaccncaac tgaagcttaa aaaatctatc acatgtataa tacctttnga agaacattaa 180
tanagcatat aaaactttta acatntgctt aatgttgtnc aattataaaa ntaatngaaa 240
aaaatgtccc tttaacatnc aatatcccac atagtgttat ttnaggggat taccnngnaa 300
naaaaaaagg gtagaaggga tttaatgaaa actctgcttn ccatttctgt ttanaaacgt 360
ctccagaaca aaaacttntc aantctttca gctaaccgca tttgagctna ggccactcaa 420
aaactccatt agncccaact tctaanggtc tctanagctt actaancctt ttgaccctt 480
accctggnta ctctgccc ca 502
```

<210> 109

<211> 1308

<212> DNA

<213> Homo sapiens

<400> 109

```
acccgaggtc tcgctaaaaat catcatggat tcacttggcg ccgtcagcac tcgacttggg 60
tttgatcttt tcaaagagct gaagaaaaca aatgatggca acatcttctt ttcccctgtg 120
ggcatcttga ctgcaatttg catggtcctc ctggggaccc gaggagccac cgcttcccag 180
ttggaggagg tgtttcactc tgaaaaagag acgaagagct caagaataaa ggctgaagaa 240
aaagaggtag ttgagaacac agaagcagta catcaacaat tccaaaagtt tttgactgaa 300
ataagcaaac tctaataatga ttatgaactg aacataacca acaggctggt tggagaaaaa 360
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<210> 110

<211> 391

<212> PRT

<213> Homo sapiens

<400> 110

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          20          25          30
Gly Ile Leu Thr Ala Ile Gly Met Val Leu Leu Gly Thr Arg Gly Ala
          35          40          45
Thr Ala Ser Gln Leu Glu Glu Val Phe His Ser Glu Lys Glu Thr Lys
          50          55          60
Ser Ser Arg Ile Lys Ala Glu Glu Lys Glu Val Ile Glu Asn Thr Glu
65          70          75          80
Ala Val His Gln Gln Phe Gln Lys Phe Leu Thr Glu Ile Ser Lys Leu
          85          90          95
Thr Asn Asp Tyr Glu Leu Asn Ile Thr Asn Arg Leu Phe Gly Glu Lys
          100          105          110
Thr Tyr Leu Phe Leu Gln Lys Tyr Leu Asp Tyr Val Glu Lys Tyr Tyr
          115          120          125
His Ala Ser Leu Glu Pro Val Asp Phe Val Asn Ala Ala Asp Glu Ser
          130          135          140
Arg Lys Lys Ile Asn Ser Trp Val Glu Ser Lys Thr Asn Glu Lys Ile
          145          150          155          160
Lys Asp Leu Phe Pro Asp Gly Ser Ile Ser Ser Ser Thr Lys Leu Val
          165          170          175
Leu Val Asn Met Val Tyr Phe Lys Gly Gln Trp Asp Arg Glu Phe Lys
          180          185          190
Lys Glu Asn Thr Lys Glu Glu Lys Phe Trp Met Asn Lys Ser Thr Ser
          195          200          205
Lys Ser Val Gln Met Met Thr Gln Ser His Ser Phe Ser Phe Thr Phe
          210          215          220
Leu Glu Asp Leu Gln Ala Lys Ile Leu Gly Ile Pro Tyr Lys Asn Asn
          225          230          235          240
Asp Leu Ser Met Phe Val Leu Leu Pro Asn Asp Ile Asp Gly Leu Glu
          245          250          255
Lys Ile Ile Asp Lys Ile Ser Pro Glu Lys Leu Val Glu Trp Thr Ser
          260          265          270
Pro Gly His Met Glu Glu Arg Lys Val Asn Leu His Leu Pro Arg Phe
          275          280          285
Glu Val Glu Asp Ser Tyr Asp Leu Glu Ala Val Leu Ala Ala Met Gly
          290          295          300

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Met Gly Asp Ala Phe Ser Glu His Lys Ala Asp Tyr Ser Gly Met Ser  
 305 310 315 320  
 Ser Gly Ser Gly Leu Tyr Ala Gln Lys Phe Leu His Ser Ser Phe Val  
 325 330 335  
 Ala Val Thr Glu Glu Gly Thr Glu Ala Ala Ala Thr Gly Ile Gly  
 340 345 350  
 Phe Thr Val Thr Ser Ala Pro Gly His Glu Asn Val His Cys Asn His  
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 Pro Phe Leu Phe Phe Ile Arg His Asn Glu Ser Asn Ser Ile Leu Phe  
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 Phe Gly Arg Phe Ser Ser Pro  
 385 390

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 <211> 1419  
 <212> DNA  
 <213> Homo sapiens

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 ggcgccgtca gcactcgact tgggtttgat cttttcaaag agctgaagaa aacaaatgat 180  
 ggcaacatct tcttttcccc tgtgggcac ttgactgcaa ttggcatggc cctcctgggg 240  
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 attgagaaca cagaagcagt acatcaacaa ttccaaaagt ttttgactga aataagcaaa 420  
 ctactaatg attatgaact gaacataacc aacaggctgt ttggagaaaa aacatacctc 480  
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 gtgctgggtga acatggttta ttttaaaggg caatgggaca gggagtttaa gaaagaaaat 720  
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 ccatataaaa acaacgacct aagcatgttt gtgcttctgc ccaacgacat cgatggcctg 900  
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 cacaatgaat ccaacagcat cctcttcttc ggcagatttt cttctcotta agatgatcgt 1320  
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<210> 112  
 <211> 400  
 <212> PRT  
 <213> Homo sapiens

<400> 112  
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 Lys Glu Leu Lys Lys Thr Asn Asp Gly Asn Ile Phe Phe Ser Pro Val

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<210> 113
<211> 957
<212> DNA
<213> Homo sapiens
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&lt;400&gt; 113

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gaaacatgag ttcttaccag cagaagcaga cctttacccc accacctcag cttcaacagc 180
agcaggtgaa acaaccacgc cagcctccac ctcaggaaat atttggtccc acaaccaagg 240
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&lt;210&gt; 114

&lt;211&gt; 161

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 114

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Gln Gln Gln Gln Val Lys Gln Pro Ser Gln Pro Pro Pro Gln Glu Ile
20          25          30
Phe Val Pro Thr Thr Lys Glu Pro Cys His Ser Lys Val Pro Gln Pro
35          40          45
Gly Asn Thr Lys Ile Pro Glu Pro Gly Cys Thr Lys Val Pro Glu Pro
50          55          60
Gly Cys Thr Lys Val Pro Glu Pro Gly Cys Thr Lys Val Pro Glu Pro
65          70          75          80
Gly Cys Thr Lys Val Pro Glu Pro Gly Cys Thr Lys Val Pro Glu Pro
85          90          95
Gly Tyr Thr Lys Val Pro Glu Pro Gly Ser Ile Lys Val Pro Asp Gln
100         105         110
Gly Phe Ile Lys Phe Pro Glu Pro Gly Ala Ile Lys Val Pro Glu Gln
115         120         125
Gly Tyr Thr Lys Val Pro Val Pro Gly Tyr Thr Lys Val Pro Glu Pro
130         135         140
Cys Pro Ser Thr Val Thr Pro Gly Pro Ala Gln Gln Lys Thr Lys Gln
145         150         155         160
Lys

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&lt;210&gt; 115

&lt;211&gt; 506

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

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 158, 195, 210, 212, 219, 226, 246, 248, 249, 255, 258, 261,  
 263, 265, 275, 304, 317, 321, 331, 337, 340, 358, 371, 377,  
 380, 396, 450, 491  
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<400> 115  
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 tatgtaaatg gtatntcatt cgctactatn antcaatng aaatanggtc tttgggttat 240  
 gaatanntng cagcncanct nanangctgt ctgtngtatt cattgtggc atagcacctc 300  
 acancattgt aacctcnatc nagtgagaca nactagnaan ttcctagtga tggctcanga 360  
 ttccaaatgg nctcatntcn aatgttttaa agttanttaa gtgtaagaaa tacagactgg 420  
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<210> 116  
 <211> 3079  
 <212> DNA  
 <213> Homo sapiens

<400> 116  
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<210> 117
<211> 6921
<212> DNA
<213> Homo sapiens

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[illegible]

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<210> 118
<211> 946
<212> DNA
<213> Homo sapiens

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<210> 119
<211> 8948
<212> DNA
<213> Homo sapiens
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| tttctcccg  | ctcctgcccc | cggcccgctg  | ccgtctccgc | gctcgcagcg  | gcctcgggag  | 180  |
| ggcccaggt  | gcgagcagcg | acctcgcgag  | ccttcgcac  | tccgcgccgg  | ttccccggcc  | 240  |
| gtccgcctat | ccttggcccc | ctccgctttc  | tccgcgcgg  | ccgcctcgc   | ttatgcctcg  | 300  |
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| ggatcaacac | tctgggcgc  | atgatccgc   | ccgagcttg  | cccgacctg   | cgctacgagg  | 420  |
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<212> DNA
<213> Homo sapiens

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510, 511, 518, 519, 539, 554, 560, 576
<223> n = A,T,C or G

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<210> 121
<211> 619
<212> DNA
<213> Homo sapiens

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```

<220>
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<223> n = A,T,C or G

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aatgaagtcc ctggtttttc atggcaactt gatcagtaaa ggattcncct ctgtttggta 540
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<210> 122
<211> 1475

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<212> DNA  
<213> Homo sapiens

<400> 122

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```

<210> 123  
<211> 2294  
<212> DNA  
<213> Homo sapiens

<400> 123

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```

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tgatttttct gatg                                     2294

```

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<210> 124
<211> 956
<212> DNA
<213> Homo sapiens

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<400> 124
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cagattgaga acctcaagga ggagctggcc tacctgaaga agaaccacga ggaggagatg 180
aacgccctgc gaggccaggt gggtggtgag atcaatgtgg agatggacgc tgccccaggc 240
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cgcagccgcc ccatctgccc cacagtctcc ggctctcca gcctcagccc cctgcttcag 900
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```

```

<210> 125
<211> 486
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 16
<223> n = A,T,C or G

```

&lt;400&gt; 125

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agcgcaggtt ttggatacta gagaaagtca tttgcttgta ctattgccat tttagaaagc 420
tctgatgtga attcaaattt tacctctgtt acttaaagcc aacaatttta aggcagtagt 480
tttact
486

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&lt;210&gt; 126

&lt;211&gt; 3552

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 126

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gctgctgacc ctctgtatct tcagtcgtgc tggatgaagcc tgcaaaaagg tgatacttaa 180
tgtaccttct aaactagagg cagacaaaat aattggcaga gttaatttgg aagagtgtct 240
caggtctgca gacctcatcc ggtcaagtga tcctgatttc agagtcttaa atgatgggtc 300
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tggactaca gtgggggtgg tttgtgccac agacagagat gaaccggaca caatgcatac 900
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agctgcaaca aaattattga gagttaatct gtgtgaatgt actcatccaa ctcatgtctg 2100
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<210> 127  
 <211> 754  
 <212> DNA  
 <213> Homo sapiens

<400> 127  
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 gctctagtgt ccatgcttct caaccattat gacccaatat tcaaccaa atcaactgaa 180  
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 cttttatggg ttgatcatct tgtcattaaa gttcaggcgt tatctatcct gtaagtggca 480  
 gaatcaagac tgcaatatcg cctgcttttc tttttaactc atgttttccc ttgactacac 540  
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 accaccttct aatactttta atacccaatc aaaatttatt atacatatgt atcatagata 660  
 ctcatctgta aagctgtgct tcaaaatagt gatctcttcc caacattaca atatatatta 720  
 atgatgtcga acctgcccgg gcggccgctc gaag 754

<210> 128  
 <211> 374  
 <212> DNA  
 <213> Homo sapiens

<400> 128  
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 ttcccttgcc cttggtaagt aactcttgat ggagaaagga ttaaagactc ttattttaacc 180

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aaaaaacaga gccagctaat catttccaaa ggtagtata tccctgctga cctcttcttt 240
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aacttaaaaa gctg                                     374

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<210> 129
<211> 546
<212> DNA
<213> Homo sapiens

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<400> 129
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aacctggtag atacatagca tgactccctg gaatagagtg ggctgggggtg cttatgctgg 240
gagagtgatt gacatgcact ttcaagctat atctaccatt tgcagcaaag gagaaaaaat 300
acctcgagta aattccatca ttttttataa catcagcacc tgctccatca tcaaggagtc 360
tcagcgtaac aggatctcca gtctctggct caactgtggc agtgacagtg gcattaagaa 420
tgggataaaa tccctgtttc acattggcat aaatcatcac aggatgagga aaatggaggc 480
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tcgaaa                                     546

```

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<210> 130
<211> 5156
<212> DNA
<213> Homo sapiens

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<210> 131
<211> 671
<212> DNA
<213> Homo sapiens

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cctgaccctt cctgctcccc aggaaggagg gtcagccccg tttgcaaaac acaggatgcc 600
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ttaactgcta t 671

```

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<210> 132
<211> 590
<212> DNA
<213> Homo sapiens

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<400> 132
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<210> 133
<211> 581
<212> DNA
<213> Homo sapiens

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ctgtcctcat ctctgcaaa ttcagcttcc ttccccaggt ctctgtgcac tctgtcttgg 180

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atgctctggg gagctcatgg gtggaggagt ctccaccaga gggagggtca ggggactggt 240
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<210> 134
<211> 4797
<212> DNA
<213> Homo sapiens

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<220>
<221> misc_feature
<222> 135, 501, 4421, 4467, 4468, 4698
<223> n = A,T,C or G

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<400> 134
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tgtggggact gggggagaga agtcccttcca tccaagctcc ctcccttact tcccccttcc 780
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aacgttatac cagtcatttt atttatagct tcgtggattt acgcttacac taaatagtct 2100

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135 501 4421 4467 4468 4698  
 n = A,T,C or G

```

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<210> 135
<211> 2856
<212> DNA
<213> Homo sapiens

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cgcacgcccc tgcgccacccg cgtacccggc gcagccagag ccaccagcgc agcgtctgcca 180
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&lt;210&gt; 136

&lt;211&gt; 356

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 136

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<210> 137
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<212> DNA
<213> Homo sapiens

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<220>
<221> misc_feature
<222> 254, 264, 279, 281, 290, 328, 342
<223> n = A,T,C or G

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<400> 137
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gtcactggct gcccccgaa cagggcgtg ctccatggct ctgcttggg tagtctgtgg 120
ctatgtctcc cagcaaggac agaaactcag aaaaatcaat cttcttacc tcattcttgt 180
cctttttctc aaagacatcg gcgaggtaat ttgtgccctt ttacctcgg ccgcgacca 240
cgctaaggcc aaanttccag acanayggcc gggccggtnc nataggggan cccaacttgg 300
ggacccaaac tctggcgcgg aaacacangg gcataagctt gnttcctgtg gggaaa 356

```

```

<210> 138
<211> 353
<212> DNA
<213> Homo sapiens

```

```

<400> 138
agggtccagtc ctccacttgg cctgatgaga gtggggagtg gcaagggacg tttctcctgc 60
aatagacact tagatttctc tcttgtggga agaaaccacc tgtccatcca ctgactcttc 120
tacattgatg tggaaattgc tgctgctacc accacctcct gaagaggctt cctgatgcc 180
aatgccagcc atcttggcat cctggccctc gagcaggetg cggtaaagtag cgatctcctg 240
ctccagccgt gtctttatgt caagcagcat ctgtactcc tggttctgag cctccatctc 300
gcatcgagac tctctcagac ctgcscgsg mssmcgctam gccgaattcc agc 353

```

```

<210> 139
<211> 371
<212> DNA
<213> Homo sapiens

```

```

<400> 139
agcgtggtcg cggccgaggt ccatccgaag caagattgca gatggcagtg tgaagagaga 60
agacatattc tacacttcaa agctttgggt caattcccat cgaccagagt tgggtccgacc 120
agccttggaa aggtcactga aaaatcttca attggattat gttgacctct acctattca 180
ttttccagtg tctgtaaagc cagggtgagga agtgatccca aaagatgaaa atggaaaaat 240
actatttgac acagtggatc tctgtgccac gtgggaggcc gtggagaagt gtaaagatgc 300
aggattggac ctgcccgggc ggccgctcga aagccgaatt ccagcacact ggcggccggt 360
actagtggat c 371

```

```

<210> 140
<211> 370
<212> DNA

```

<213> Homo sapiens

<400> 140

```
tagcgtggtc gcggccgaggg tccatctccc tttgggaact agggggctgc tgggtgggaaa 60
tgggagccag ggcagatggt gcatttccttt gtgtccctgt aaatgtggga ctacaagaag 120
aggagctgcc tgagtgggtac tttctcttcc tggtaatcct ctggcccagc ctcatggcag 180
aatagaggta tttttaggct atttttgtaa tatggcttct ggtcaaaatc cctgtgtagc 240
tgaattccca agccctgcat tgtacagccc cccactcccc tcaccaccta ataaaggaat 300
agttaacact caaaaaaaaa aaaaaaacctg cccgggcggc cgctcgaaag ccgaattcca 360
gcacactggc                                     370
```

<210> 141

<211> 371

<212> DNA

<213> Homo sapiens

<400> 141

```
tagcgtggtc gcggccgaggg tcctctgtgc tgctgtcac agcccgatgg taccagcgca 60
gggtgtaggc agtgcaggag cctcatcca gtggcaggga acaggggtca tcaactatccc 120
aaggagcttc agggctcctg tactcctcca cagaatactc ggagtattca gactactcat 180
catcctcagg gggtaaccgc tcttctcct ctgcatgaga gacgcggagc acaggcacag 240
catggagctg ggagccggca gtgtctgcag cataactagg gaggggtcgt gatccagatg 300
cgatgaactg gccctggcag gcacagtgtc gactcatctc ttggcgacct gcccgggcgg 360
ccgctcgaag c                                     371
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<210> 142

<211> 343

<212> DNA

<213> Homo sapiens

<400> 142

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gcgttttgag gccaatggtg taaaaggaaa tatcttcaca taaaaactag atggaagcat 60
tgtcagaaac ctctttgtga tgtttgcttt caactcacag agttgaacat tccttttcat 120
agagcagttt tgaacactc tttttagtaa tttgcaagcg gatgattgga tcgctatgag 180
gtcttcattg gaaacgggat acctttacat aaaaactaga cagtagcatt ctcagaaatt 240
tctttgggat gtgggcattc aaccacaga ggagaacttc atttgataga gcagttttga 300
aacacccttt ttgtagaatc tacaggtgga catttagagt gct                                     343
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<210> 143

<211> 354

<212> DNA

<213> Homo sapiens

<400> 143

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aggctctgat gcagaaaaac tcagactgtc tgcaacttta cagatggtgc attggttcag 60
catcaggagt gggatgggaa ggaaagcaca ataacaagaa aattgaaaga tgggaaatta 120
gtggtggagt gtgtcatgaa caatgtcacc tgtactcgga tctatgaaaa agtagaataa 180
aaattccatc atcactttgg acaggagtta attaagagaa tgaccaagct cagttcaatg 240
agcaaatctc catactgttt ctttcttttt tttttcatta ctgtgttcaa ttatctttat 300
cataaacatt ttacatgcag ctatttcaaa gtgtgttgga ttaattagga tcat                                     354
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<210> 144

<211> 353

<212> DNA

<213> Homo sapiens

<400> 144

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cctagagcac atctggatct cagccccacc cctggcaacc tgccctgccta gagaactccc 120
aagatgacag actaagtagg attctgccat ttagaataat tctggtatcc tgggcgttgc 180
gttaagtgtc ttaactttca ttctgtctta cgatagtctt cagaggtggg aacagatgaa 240
gaaaccatgc cccagagaag gttaagtgc ttctcttcta tggagccagt gttccaacct 300
aggtttgcct gataccagac ctgtggcccc acctcccatg caggtctctg tgg          353

```

<210> 145

<211> 371

<212> DNA

<213> Homo sapiens

<400> 145

```

caggtctgtc ataaactggg ctggagtttc tgacgactcc ttgttcacca aatgcacat 60
ttcctgagac ttgttgacct ctccgttgag tccacttggc tttctgtcct ccacagctcc 120
attgccactg ttgatcacta gctttttctt ctgccacac cttcttcgac tgttgactgc 180
aatgcaaaact gcaagaatca aagccaaggc caagagggat gccaagatga tcagccattc 240
tggaatttgg ggtgtcctta taggaccaga ggttgtgttt gctccacctt cttgactccc 300
atgtgagacc tcggcccgca ccacgctaag ccgaattcca gcacactggc ggcccgttac 360
tagtggatcc g          371

```

<210> 146

<211> 355

<212> DNA

<213> Homo sapiens

<400> 146

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caggatggcg agtagcagcg gctccaaggc tgaattcatt gtccggagga aatataaact 120
ggtacggaag atcgggtctg gctccttcgg ggacatctat ttggcgatca acatcaccaa 180
cggcgaggaa gtggcagtga agctagaatc tcagaaggcc aggcacccc agttgctgta 240
cgagagcaag ctctataaga ttcttcaagg tggggttggc atccccaca tacggtggta 300
tggtcaggaa aaagactaca atgtactagt catggatctt ctgggacctc gcctc          355

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<210> 147

<211> 355

<212> DNA

<213> Homo sapiens

<400> 147

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tactatgcac gtgctgtgat ttgaaacata actcgtccca aaaacttgct acgatcatcc 120
tgacttttta ggttggctga tccatcaatc ttgcaactca ctgttacttc tttcccagtg 180
ttgttaggag caaagctgac ctgaacagca accaatggct gtagataccc aacatgcagt 240
tttttcccat aatatgggaa atattttaag tctatcattc cattatgagg ataaactgct 300
acatttggtg tatcttcatt ctttgaacaa caatctatcc ttggcactcc ttcag          355

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<210> 148

<211> 369

<212> DNA

<213> Homo sapiens

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<400> 148
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agggagtgtg ccgagggctt ctgagaaggt ttctctcaca tctagaaaga agcgcttaag 180
atgtggcagc cctctctctt caagtggctc ttgtcctgtt gccctgggag ttctcaaatt 240
gctgcagcag cctccatcca gcctgaggat gacatcaata cacagaggaa gaagagtcag 300
gaaaagatga gagaagttac agactctcct gggcgacccc gagagcttac cattcctcag 360
acttcttca
369

```

```

<210> 149
<211> 620
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 169, 171, 222, 472, 528, 559, 599
<223> n = A,T,C or G

```

```

<400> 149
actagtcaaa aatgctaaaa taatttggga gaaaatattt ttttaagtagt gttatagttt 60
catgtttatc ttttattatg ttttgtgaag ttgtgtcttt tcactaatta cctatactat 120
gccaatattt ccttataatc atccataaca tttatactac atttghtaana naatatgcac 180
gtgaaactta acactttata aggtaaaaat gaggtttcca anatttaata atctgatcaa 240
gttcttgtaa tttccaaata gaatggactt ggtctgttaa gggctaagga gaagaggaag 300
ataagggttaa aagttgttaa tgaccaaaca ttctaaaaga aatgcaaaaa aaaagtattt 360
tttcaagcct tcgaactatt taaggaaaagc aaaatcattt cctaaatgca tatcatttgt 420
gagaatttct cattaatatc ctgaatcatt catttacta aggctcatgt tnactccgat 480
atgtctctaa gaaagtacta tttcatggtc caaacctggg tgccatantt gggtaaaggc 540
tttcccttaa gtgtgaaant atttaaaatg aaattttcct ctttttaaaa attctttana 600
agggttaagg gtgttgggga
620

```

```

<210> 150
<211> 371
<212> DNA
<213> Homo sapiens

```

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<400> 150
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gagcaaccag tatcacttcc ctgtttataa aacctctaac catctctttg ttctttgaac 120
atgctgaaaa ccacctgggc tgcattgtatg cccgaatttg yaattctttt ctctcaaattg 180
aaaatttaaa ttttagggatt catttctata ttttcacata tgtagtatta ttatttcctt 240
atatgtgtaa ggtgaaattt atggattttg agtgtgcaag aaaatatatt tttaaagctt 300
tcatttttcc ccagtgaaat gatttagaat tttttatgta aatatacaga atgttttttc 360
ttacttttat a
371

```

```

<210> 151
<211> 4655
<212> DNA
<213> Homo sapiens

```

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<400> 151
gggacttgag ttctgttata ttcttaagta gattcatatt gtaagggtct cgggggtgggg 60

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| gggttggcaa  | aatcctggag  | ccagaagaaa  | ggacagcagc  | attgatcaat  | cttacagcta  | 120  |
| acatgttgta  | cctggaaaac  | aatgcccgaga | ctcaatttag  | tgagccacag  | tacacgaacc  | 180  |
| tggggctcct  | gaacagcatg  | gaccagcaga  | ttcagaacgg  | ctcctcgtec  | accagtcctt  | 240  |
| ataacacaga  | ccacgcgcag  | aacagcgtca  | cgggcgccctc | gccctaagca  | cagcccagct  | 300  |
| ccaccttcga  | tgtctctctt  | ccatcaccgc  | ccatcccctc  | caacaccgac  | taccagggcc  | 360  |
| cgcacagttt  | cgacgtgtcc  | ttccagcagt  | cgagcacccg  | caagtgcggc  | acctggagct  | 420  |
| attccactga  | actgaagaaa  | ctctactgcc  | aaattgcaaa  | gacatgcccc  | atccagatca  | 480  |
| aggtgatgac  | cccacctctt  | cagggagctg  | ttatccgcgc  | catgcctgtc  | tacaaaaaag  | 540  |
| ctgagcacgt  | cacggagggtg | gtgaagcgtg  | gccccaaaca  | tgagctgagc  | cgtaaaattca | 600  |
| acgagggaca  | gattggccctt | yctagtcttt  | tgatttcgagt | agaggggaaac | agccatgccc  | 660  |
| agtatgtaga  | agatcccctc  | acaggaagac  | agagtgtgct  | ggtaccttat  | gagccacccc  | 720  |
| aggttggcac  | tgaattcacg  | acagtcttgt  | acaatttcat  | gtgtaacagc  | agttgtgttg  | 780  |
| gagggatgaa  | ccgcgcgtcca | attttaatat  | ttgttactct  | ggaaaccaga  | gatgggcaag  | 840  |
| tcttgggccc  | acgtctgttt  | gaggcccgga  | tctgtgcttg  | cccaggaaga  | gacaggaagg  | 900  |
| cggtatgaaga | tagcatcaga  | aagcagcaag  | tttcggacag  | tacaaagaac  | ggtgatggta  | 960  |
| cgaagcgcgc  | gtttcgctcag | aacacacatg  | gtatccagat  | gacatccctc  | aagaaacgaa  | 1020 |
| gatccccaga  | tgatgaactg  | gtatacttac  | cagtgagggg  | ccgtgagact  | tatgaaatgc  | 1080 |
| tggtgaagat  | caaagagtc   | ctggaactca  | tgcagtacct  | tcttcagcac  | acaattgaaa  | 1140 |
| cgtacaggga  | acagcaacag  | cagcagcacc  | agcacttact  | tcagaaacag  | acctcaatac  | 1200 |
| agtctccatc  | ttcatatggt  | aacagctccc  | cacctctgaa  | caaaatgaac  | agcatgaaca  | 1260 |
| agctgccttc  | tgtgagccag  | cttatcaacc  | ctcagcagcg  | caacgcctc   | actcctacaa  | 1320 |
| ccattcctga  | tggcatggga  | gccaacattc  | ccatgatggg  | caccacatg   | ccaatggctg  | 1380 |
| gagacatgaa  | tggactcagc  | cccacccagg  | cactccctcc  | cccactctcc  | atgccatcca  | 1440 |
| cctcccactg  | cacaccccca  | cctccgtatc  | ccacagattg  | cagcattgtc  | agtttcttag  | 1500 |
| cgaggttggg  | ctgttcatca  | tgtctggact  | atttcacgac  | ccaggggctg  | accaccatct  | 1560 |
| atcagattga  | gcattactcc  | atggatgac   | tggcaagtct  | gaaaatccct  | gagcaatttc  | 1620 |
| gacatgcgat  | ctggaagggc  | atcctggacc  | accggcagct  | ccacgaattc  | tcttcccctt  | 1680 |
| ctcatctcct  | gcggacccca  | agcagtgcct  | ctacagtcag  | tgtgggctcc  | agtgagaccc  | 1740 |
| ggggtgagcg  | tgttattgat  | gctgtgcgat  | tcacctccg   | ccagaccatc  | tctttcccac  | 1800 |
| cccagagatga | gtggaatgac  | ttcaactttg  | acatggatgc  | tgcgcgcaat  | aagcaacagc  | 1860 |
| gcatcaaaaga | ggagggggag  | tgagcctcac  | catgtgagct  | cttctctacc  | ctctcctaac  | 1920 |
| tgcagccccc  | ctaaaagcac  | tcttgcttaa  | tcttcaaagc  | cttctcccta  | gtcctccccc  | 1980 |
| ttctcttgt   | ctgatttctt  | aggggaagga  | gaagtaagag  | gcttacttct  | taccttaacc  | 2040 |
| atctgacctg  | gcatctaatt  | ctgattctgg  | ctttaagcct  | tcaaaactat  | agcttgacga  | 2100 |
| actgtagctt  | gccatggcta  | ggtagaagtg  | agcaaaaaag  | agttgggtgt  | ctccttaagc  | 2160 |
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| atataaatgt  | ataaatatac  | agtatagatt  | tttgggtggg  | gggcattgag  | tattgtttta  | 2280 |
| aatgtaat    | aaatgaaaga  | aaattgagtt  | gcacttattg  | accatttttt  | aatttacttg  | 2340 |
| ttttggatgg  | cttgtctata  | ctccttccct  | taaggggtat  | catgtatggt  | gatagggtatc | 2400 |
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| aagactgtag  | atatgtattc  | ttttctcagt  | gttggtatat  | tttatattac  | tgacatttct  | 2580 |
| tctagtgtg   | atggttcacg  | ttggggtgat  | ttaatccagt  | tataagaaga  | agttcatgtc  | 2640 |
| caaacgtcct  | ctttagtttt  | tgggttggga  | tgaggaaaaat | tcttaaaagg  | cccatagcag  | 2700 |
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| taccagatac  | cttatcttac  | aatattgatt  | gggaaaacat  | ttgctgccat  | tacagaggta  | 2820 |
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| gaattctgat  | tgatttgatt  | gggatgaatg  | ccatctatct  | agttctaaca  | gtgaagtttt  | 2940 |
| actgtctatt  | aatattcagg  | gtaaaatagga | atcattcaga  | aatgttgagt  | ctgtactaaa  | 3000 |
| cagtaagata  | tctcaatgaa  | ccataaatcc  | aactttgtaa  | aaatcttttg  | aagcatagat  | 3060 |
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| agtggtgatt  | tgaaaaatat  | aaaattatga  | gattgggttt  | cctgtggcat  | aaattgcac   | 3240 |
| actgtatcat  | tttctttttt  | aaccgggtaag | agtttcagtt  | tgttggaagg  | taactgtgag  | 3300 |

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```

<210> 152

<211> 586

<212> PRT

<213> Homo sapiens

<400> 152

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Met Leu Tyr Leu Glu Asn Asn Ala Gln Thr Gln Phe Ser Glu Pro Gln
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Tyr Thr Asn Leu Gly Leu Leu Asn Ser Met Asp Gln Gln Ile Gln Asn
      20             25             30
Gly Ser Ser Ser Thr Ser Pro Tyr Asn Thr Asp His Ala Gln Asn Ser
      35             40             45
Val Thr Ala Pro Ser Pro Tyr Ala Gln Pro Ser Ser Thr Phe Asp Ala
      50             55             60
Leu Ser Pro Ser Pro Ala Ile Pro Ser Asn Thr Asp Tyr Pro Gly Pro
      65             70             75             80
His Ser Phe Asp Val Ser Phe Gln Gln Ser Ser Thr Ala Lys Ser Ala
      85             90             95
Thr Trp Thr Tyr Ser Thr Glu Leu Lys Lys Leu Tyr Cys Gln Ile Ala
      100            105            110
Lys Thr Cys Pro Ile Gln Ile Lys Val Met Thr Pro Pro Pro Gln Gly
      115            120            125
Ala Val Ile Arg Ala Met Pro Val Tyr Lys Lys Ala Glu His Val Thr
      130            135            140
Glu Val Val Lys Arg Cys Pro Asn His Glu Leu Ser Arg Glu Phe Asn
      145            150            155            160
Glu Gly Gln Ile Ala Pro Ser Ser His Leu Ile Arg Val Glu Gly Asn
      165            170            175
Ser His Ala Gln Tyr Val Glu Asp Pro Ile Thr Gly Arg Gln Ser Val
      180            185            190

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Leu Val Pro Tyr Glu Pro Pro Gln Val Gly Thr Glu Phe Thr Thr Val  
 195 200 205  
 Leu Tyr Asn Phe Met Cys Asn Ser Ser Cys Val Gly Gly Met Asn Arg  
 210 215 220  
 Arg Pro Ile Leu Ile Ile Val Thr Leu Glu Thr Arg Asp Gly Gln Val  
 225 230 235 240  
 Leu Gly Arg Arg Cys Phe Glu Ala Arg Ile Cys Ala Cys Pro Gly Arg  
 245 250 255  
 Asp Arg Lys Ala Asp Glu Asp Ser Ile Arg Lys Gln Gln Val Ser Asp  
 260 265 270  
 Ser Thr Lys Asn Gly Asp Gly Thr Lys Arg Pro Phe Arg Gln Asn Thr  
 275 280 285  
 His Gly Ile Gln Met Thr Ser Ile Lys Lys Arg Arg Ser Pro Asp Asp  
 290 295 300  
 Glu Leu Val Tyr Leu Pro Val Arg Gly Arg Glu Thr Tyr Glu Met Leu  
 305 310 315 320  
 Val Lys Ile Lys Glu Ser Leu Glu Leu Met Gln Tyr Leu Leu Gln His  
 325 330 335  
 Thr Ile Glu Thr Tyr Arg Gln Gln Gln Gln Gln His Gln His Leu  
 340 345 350  
 Leu Gln Lys Gln Thr Ser Ile Gln Ser Pro Ser Ser Tyr Gly Asn Ser  
 355 360 365  
 Ser Pro Pro Leu Asn Lys Met Asn Ser Met Asn Lys Leu Pro Ser Val  
 370 375 380  
 Ser Gln Leu Ile Asn Pro Gln Gln Arg Asn Ala Leu Thr Pro Thr Thr  
 385 390 395 400  
 Ile Pro Asp Gly Met Gly Ala Asn Ile Pro Met Met Gly Thr His Met  
 405 410 415  
 Pro Met Ala Gly Asp Met Asn Gly Leu Ser Pro Thr Gln Ala Leu Pro  
 420 425 430  
 Pro Pro Leu Ser Met Pro Ser Thr Ser His Cys Thr Pro Pro Pro Pro  
 435 440 445  
 Tyr Pro Thr Asp Cys Ser Ile Val Ser Phe Leu Ala Arg Leu Gly Cys  
 450 455 460  
 Ser Ser Cys Leu Asp Tyr Phe Thr Thr Gln Gly Leu Thr Thr Ile Tyr  
 465 470 475 480  
 Gln Ile Glu His Tyr Ser Met Asp Asp Leu Ala Ser Leu Lys Ile Pro  
 485 490 495  
 Glu Gln Phe Arg His Ala Ile Trp Lys Gly Ile Leu Asp His Arg Gln  
 500 505 510  
 Leu His Glu Phe Ser Ser Pro Ser His Leu Leu Arg Thr Pro Ser Ser  
 515 520 525  
 Ala Ser Thr Val Ser Val Gly Ser Ser Glu Thr Arg Gly Glu Arg Val  
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 Ile Asp Ala Val Arg Phe Thr Leu Arg Gln Thr Ile Ser Phe Pro Pro  
 545 550 555 560  
 Arg Asp Glu Trp Asn Asp Phe Asn Phe Asp Met Asp Ala Arg Arg Asn  
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&lt;210&gt; 153

&lt;211&gt; 2007



<212> DNA  
<213> Homo sapiens

<400> 153

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gttgattgac taaaaaaaaa aaaaaaa 2007

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<210> 154  
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<212> DNA  
<213> Homo sapiens

<400> 154

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<210> 155

<211> 153

<212> PRT

<213> Homo sapiens

<400> 155

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Met Thr Ser Val Arg Val Ala Ala Tyr Phe Glu Asn Phe Leu Ala Ala
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 20            25            30
Pro Met Gly Asp Val Pro Met Asp Gly Ile Ser Val Ala Asp Ile Gly
 35            40            45
Ala Ala Val Ser Ser Ile Phe Asn Ser Pro Glu Glu Phe Leu Gly Lys
 50            55            60
Ala Val Gly Leu Ser Ala Glu Ala Leu Thr Ile Gln Gln Tyr Ala Asp
 65            70            75            80
Val Leu Ser Lys Ala Leu Gly Lys Glu Val Arg Asp Ala Lys Ile Thr
 85            90            95
Pro Glu Ala Phe Glu Lys Leu Gly Phe Pro Ala Ala Lys Glu Ile Ala
100           105           110
Asn Met Cys Arg Phe Tyr Glu Met Lys Pro Asp Arg Asp Val Asn Leu
115           120           125
Thr His Gln Leu Asn Pro Lys Val Lys Ser Phe Ser Gln Phe Ile Ser
130           135           140
Glu Asn Gln Gly Ala Phe Lys Gly Met
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<210> 156  
 <211> 128  
 <212> PRT  
 <213> Homo sapiens

<400> 156  
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 20 25 30  
 Pro Met Gly Asp Val Pro Met Asp Gly Ile Ser Val Ala Asp Ile Gly  
 35 40 45  
 Ala Ala Val Ser Ser Ile Phe Asn Ser Pro Glu Glu Phe Leu Gly Lys  
 50 55 60  
 Ala Val Gly Leu Ser Ala Glu Ala Leu Thr Ile Gln Gln Tyr Ala Asp  
 65 70 75 80  
 Val Leu Ser Lys Ala Leu Gly Lys Glu Val Arg Asp Ala Lys Thr Ile  
 85 90 95  
 Cys Ala Ile Asp Asp Gln Lys Thr Val Glu Glu Gly Phe Met Glu Asp  
 100 105 110  
 Val Gly Leu Ser Trp Ser Leu Arg Glu His Asp His Val Ala Gly Ala  
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<210> 157  
 <211> 424  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
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<400> 157  
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<210> 158  
 <211> 2099  
 <212> DNA  
 <213> Homo sapiens

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<210> 159

<211> 291

<212> PRT

<213> Homo sapiens

<400> 159

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 20          25          30
Val Met Ile Leu Val Val Ala Ala Gln Glu Val Trp Gly Asp Glu Gln
 35          40          45
Glu Asp Phe Val Cys Asn Thr Leu Gln Pro Gly Cys Lys Asn Val Cys
 50          55          60
Tyr Asp His Phe Phe Pro Val Ser His Ile Arg Leu Trp Ala Leu Gln
 65          70          75          80
Leu Ile Phe Val Ser Thr Pro Ala Leu Leu Val Ala Met His Val Ala
 85          90          95
Tyr Tyr Arg His Glu Thr Thr Arg Lys Phe Arg Arg Gly Glu Lys Arg
100          105          110

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Asn Asp Phe Lys Asp Ile Glu Asp Ile Lys Lys Gln Lys Val Arg Ile  
 115 120 125  
 Glu Gly Ser Leu Trp Trp Thr Tyr Thr Ser Ser Ile Phe Phe Arg Ile  
 130 135 140  
 Ile Phe Glu Ala Ala Phe Met Tyr Val Phe Tyr Phe Leu Tyr Asn Gly  
 145 150 155 160  
 Tyr His Leu Pro Trp Val Leu Lys Cys Gly Ile Asp Pro Cys Pro Asn  
 165 170 175  
 Leu Val Asp Cys Phe Ile Ser Arg Pro Thr Glu Lys Thr Val Phe Thr  
 180 185 190  
 Ile Phe Met Ile Ser Ala Ser Val Ile Cys Met Leu Leu Asn Val Ala  
 195 200 205  
 Glu Leu Cys Tyr Leu Leu Leu Lys Val Cys Phe Arg Arg Ser Lys Arg  
 210 215 220  
 Ala Gln Thr Gln Lys Asn His Pro Asn His Ala Leu Lys Glu Ser Lys  
 225 230 235 240  
 Gln Asn Glu Met Asn Glu Leu Ile Ser Asp Ser Gly Gln Asn Ala Ile  
 245 250 255  
 Thr Gly Ser Gln Ala Lys His Phe Lys Val Lys Cys Ser Cys Val Ile  
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 Ser Val Ala  
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<210> 160

<211> 3951

<212> DNA

<213> Homo sapiens

<400> 160

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<210> 161
<211> 943
<212> PRT
<213> Homo sapiens

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<400> 161
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1           5           10           15
Thr Leu Leu Val Ala Leu Ser Ser Glu Leu Pro Phe Leu Gly Ala Gly

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|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
|     |     |     | 20  |     |     |     |     | 25  |     |     |     |     | 30  |     |     |
| Val | Gln | Leu | Gln | Asp | Asn | Gly | Tyr | Asn | Gly | Leu | Leu | Ile | Ala | Ile | Asn |
|     |     | 35  |     |     |     |     | 40  |     |     |     |     | 45  |     |     |     |
| Pro | Gln | Val | Pro | Glu | Asn | Gln | Asn | Leu | Ile | Ser | Asn | Ile | Lys | Glu | Met |
|     | 50  |     |     |     |     | 55  |     |     |     |     | 60  |     |     |     |     |
| Ile | Thr | Glu | Ala | Ser | Phe | Tyr | Leu | Phe | Asn | Ala | Thr | Lys | Arg | Arg | Val |
| 65  |     |     |     |     | 70  |     |     |     |     | 75  |     |     |     |     | 80  |
| Phe | Phe | Arg | Asn | Ile | Lys | Ile | Leu | Ile | Pro | Ala | Thr | Trp | Lys | Ala | Asn |
|     |     |     |     | 85  |     |     |     |     | 90  |     |     |     |     | 95  |     |
| Asn | Asn | Ser | Lys | Ile | Lys | Gln | Glu | Ser | Tyr | Glu | Lys | Ala | Asn | Val | Ile |
|     |     |     | 100 |     |     |     |     | 105 |     |     |     |     | 110 |     |     |
| Val | Thr | Asp | Trp | Tyr | Gly | Ala | His | Gly | Asp | Asp | Pro | Tyr | Thr | Leu | Gln |
|     |     | 115 |     |     |     |     | 120 |     |     |     |     | 125 |     |     |     |
| Tyr | Arg | Gly | Cys | Gly | Lys | Glu | Gly | Lys | Tyr | Ile | His | Phe | Thr | Pro | Asn |
|     | 130 |     |     |     |     | 135 |     |     |     |     | 140 |     |     |     |     |
| Phe | Leu | Leu | Asn | Asp | Asn | Leu | Thr | Ala | Gly | Tyr | Gly | Ser | Arg | Gly | Arg |
| 145 |     |     |     |     | 150 |     |     |     |     | 155 |     |     |     |     | 160 |
| Val | Phe | Val | His | Glu | Trp | Ala | His | Leu | Arg | Trp | Gly | Val | Phe | Asp | Glu |
|     |     |     |     | 165 |     |     |     |     | 170 |     |     |     |     | 175 |     |
| Tyr | Asn | Asn | Asp | Lys | Pro | Phe | Tyr | Ile | Asn | Gly | Gln | Asn | Gln | Ile | Lys |
|     |     |     | 180 |     |     |     |     | 185 |     |     |     |     | 190 |     |     |
| Val | Thr | Arg | Cys | Ser | Ser | Asp | Ile | Thr | Gly | Ile | Phe | Val | Cys | Glu | Lys |
|     |     | 195 |     |     |     |     | 200 |     |     |     |     | 205 |     |     |     |
| Gly | Pro | Cys | Pro | Gln | Glu | Asn | Cys | Ile | Ile | Ser | Lys | Leu | Phe | Lys | Glu |
|     | 210 |     |     |     |     | 215 |     |     |     |     | 220 |     |     |     |     |
| Gly | Cys | Thr | Phe | Ile | Tyr | Asn | Ser | Thr | Gln | Asn | Ala | Thr | Ala | Ser | Ile |
| 225 |     |     |     |     | 230 |     |     |     |     | 235 |     |     |     |     | 240 |
| Met | Phe | Met | Gln | Ser | Leu | Ser | Ser | Val | Val | Glu | Phe | Cys | Asn | Ala | Ser |
|     |     |     |     | 245 |     |     |     |     | 250 |     |     |     |     | 255 |     |
| Thr | His | Asn | Gln | Glu | Ala | Pro | Asn | Leu | Gln | Asn | Gln | Met | Cys | Ser | Leu |
|     |     |     | 260 |     |     |     |     | 265 |     |     |     |     | 270 |     |     |
| Arg | Ser | Ala | Trp | Asp | Val | Ile | Thr | Asp | Ser | Ala | Asp | Phe | His | His | Ser |
|     |     | 275 |     |     |     |     | 280 |     |     |     |     | 285 |     |     |     |
| Phe | Pro | Met | Asn | Gly | Thr | Glu | Leu | Pro | Pro | Pro | Pro | Thr | Phe | Ser | Leu |
|     | 290 |     |     |     |     | 295 |     |     |     |     | 300 |     |     |     |     |
| Val | Glu | Ala | Gly | Asp | Lys | Val | Val | Cys | Leu | Val | Leu | Asp | Val | Ser | Ser |
| 305 |     |     |     |     | 310 |     |     |     |     | 315 |     |     |     |     | 320 |
| Lys | Met | Ala | Glu | Ala | Asp | Arg | Leu | Leu | Gln | Leu | Gln | Gln | Ala | Ala | Glu |
|     |     |     |     | 325 |     |     |     |     | 330 |     |     |     |     | 335 |     |
| Phe | Tyr | Leu | Met | Gln | Ile | Val | Glu | Ile | His | Thr | Phe | Val | Gly | Ile | Ala |
|     |     |     | 340 |     |     |     |     | 345 |     |     |     |     | 350 |     |     |
| Ser | Phe | Asp | Ser | Lys | Gly | Glu | Ile | Arg | Ala | Gln | Leu | His | Gln | Ile | Asn |
|     |     | 355 |     |     |     |     | 360 |     |     |     |     | 365 |     |     |     |
| Ser | Asn | Asp | Asp | Arg | Lys | Leu | Leu | Val | Ser | Tyr | Leu | Pro | Thr | Thr | Val |
|     | 370 |     |     |     |     | 37  |     |     |     |     |     |     |     |     |     |

|   |     |     |     |     |
|---|-----|-----|-----|-----|
| 450   |     | 455 |     | 460 |
| Phe Phe Val Pro Asp Ile Ser Asn Ser Asn Ser Met Ile Asp Ala Phe |     |     |     |     |
| 465   |     | 470 |     | 475 |
| Ser Arg Ile Ser Ser Gly Thr Gly Asp Ile Phe Gln Gln His Ile Gln |     |     |     |     |
|   | 485 |     | 490 | 495 |
| Leu Glu Ser Thr Gly Glu Asn Val Lys Pro His His Gln Leu Lys Asn |     |     |     |     |
|   | 500 |     | 505 | 510 |
| Thr Val Thr Val Asp Asn Thr Val Gly Asn Asp Thr Met Phe Leu Val |     |     |     |     |
|   | 515 |     | 520 | 525 |
| Thr Trp Gln Ala Ser Gly Pro Pro Glu Ile Ile Leu Phe Asp Pro Asp |     |     |     |     |
|   | 530 |     | 535 | 540 |
| Gly Arg Lys Tyr Tyr Thr Asn Asn Phe Ile Thr Asn Leu Thr Phe Arg |     |     |     |     |
| 545   |     | 550 |     | 555 |
| Thr Ala Ser Leu Trp Ile Pro Gly Thr Ala Lys Pro Gly His Trp Thr |     |     |     |     |
|   | 565 |     | 570 | 575 |
| Tyr Thr Leu Asn Asn Thr His His Ser Leu Gln Ala Leu Lys Val Thr |     |     |     |     |
|   | 580 |     | 585 | 590 |
| Val Thr Ser Arg Ala Ser Asn Ser Ala Val Pro Pro Ala Thr Val Glu |     |     |     |     |
|   | 595 |     | 600 | 605 |
| Ala Phe Val Glu Arg Asp Ser Leu His Phe Pro His Pro Val Met Ile |     |     |     |     |
|   | 610 |     | 615 | 620 |
| Tyr Ala Asn Val Lys Gln Gly Phe Tyr Pro Ile Leu Asn Ala Thr Val |     |     |     |     |
| 625   |     | 630 |     | 635 |
| Thr Ala Thr Val Glu Pro Glu Thr Gly Asp Pro Val Thr Leu Arg Leu |     |     |     |     |
|   | 645 |     | 650 | 655 |
| Leu Asp Asp Gly Ala Gly Ala Asp Val Ile Lys Asn Asp Gly Ile Tyr |     |     |     |     |
|   | 660 |     | 665 | 670 |
| Ser Arg Tyr Phe Phe Ser Phe Ala Ala Asn Gly Arg Tyr Ser Leu Lys |     |     |     |     |
|   | 675 |     | 680 | 685 |
| Val His Val Asn His Ser Pro Ser Ile Ser Thr Pro Ala His Ser Ile |     |     |     |     |
|   | 690 |     | 695 | 700 |
| Pro Gly Ser His Ala Met Tyr Val Pro Gly Tyr Thr Ala Asn Gly Asn |     |     |     |     |
| 705   |     | 710 |     | 715 |
| Ile Gln Met Asn Ala Pro Arg Lys Ser Val Gly Arg Asn Glu Glu Glu |     |     |     |     |
|   | 725 |     | 730 | 735 |
| Arg Lys Trp Gly Phe Ser Arg Val Ser Ser Gly Gly Ser Phe Ser Val |     |     |     |     |
|   | 740 |     | 745 | 750 |
| Leu Gly Val Pro Ala Gly Pro His Pro Asp Val Phe Pro Pro Cys Lys |     |     |     |     |
|   | 755 |     | 760 | 765 |
| Ile Ile Asp Leu Glu Ala Val Lys Val Glu Glu Glu Leu Thr Leu Ser |     |     |     |     |
|   | 770 |     | 775 | 780 |
| Trp Thr Ala Pro Gly Glu Asp Phe Asp Gln Gly Gln Ala Thr Ser Tyr |     |     |     |     |
| 785   |     | 790 |     | 795 |
| Glu Ile Arg Met Ser Lys Ser Leu Gln Asn Ile Gln Asp Asp Phe Asn |     |     |     |     |
|   | 805 |     | 810 | 815 |
| Asn Ala Ile Leu Val Asn Thr Ser Lys Arg Asn Pro Gln Gln Ala Gly |     |     |     |     |
|   | 820 |     | 825 | 830 |
| Ile Arg Glu Ile Phe Thr Phe Ser Pro Gln Ile Ser Thr Asn Gly Pro |     |     |     |     |
|   | 835 |     | 840 | 845 |
| Glu His Gln Pro Asn Gly Glu Thr His Glu Ser His Arg Ile Tyr Val |     |     |     |     |
|   | 850 |     | 855 | 860 |
| Ala Ile Arg Ala Met Asp Arg Asn Ser Leu Gln Ser Ala Val Ser Asn |     |     |     |     |
| 865   |     | 870 |     | 875 |
| Ile Ala Gln Ala Pro Leu Phe Ile Pro Pro Asn Ser Asp Pro Val Pro |     |     |     |     |



|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
|     |     |     |     | 885 |     |     |     | 890 |     |     |     | 895 |     |     |     |
| Ala | Arg | Asp | Tyr | Leu | Ile | Leu | Lys | Gly | Val | Leu | Thr | Ala | Met | Gly | Leu |
| 900 |     |     |     |     |     |     |     | 905 |     |     |     | 910 |     |     |     |
| Ile | Gly | Ile | Ile | Cys | Leu | Ile | Ile | Val | Val | Thr | His | His | Thr | Leu | Ser |
| 915 |     |     |     |     |     |     |     | 920 |     |     |     | 925 |     |     |     |
| Arg | Lys | Lys | Arg | Ala | Asp | Lys | Lys | Glu | Asn | Gly | Thr | Lys | Leu | Leu |     |
| 930 |     |     |     | 935 |     |     |     | 940 |     |     |     |     |     |     |     |

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<210> 162
<211> 498
<212> DNA
<213> Homo sapiens
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| tggagaacca | cgtggacagc | accatgaaca | tgttggggcg | gggaggcagt | gctggccgga | 60  |
| agccctcaa  | gtcgggtatg | aaggagctgg | ccgtgttccg | ggagaaggtc | actgagcagc | 120 |
| accggcagat | gggcaagggt | ggcaagcatc | accttggcct | ggaggagccc | aagaagctgc | 180 |
| gaccacccc  | tgccaggact | ccctgccaac | aggaactgga | ccaggctctg | gagcggatct | 240 |
| ccaccatgcg | ccttccggat | gagcggggcc | ctctggagca | cctctactcc | ctgcacatcc | 300 |
| ccaactgtga | caagcatggc | ctgtacaacc | tcaaacagtg | gcaagatgtc | tctgaacggg | 360 |
| cagcgtgggg | agtgtggtg  | tgtgaacccc | aacaccggga | agctgatcca | gggagccccc | 420 |
| accatccggg | gggacccga  | gtgtcatctc | ttctacaatg | agcagcagga | ggctcgcggg | 480 |
| gtgcacaccc | cagcggat   |            |            |            |            | 498 |

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<210> 163
<211> 1128
<212> DNA
<213> Homo sapiens
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|            |            |             |            |             |             |      |
|------------|------------|-------------|------------|-------------|-------------|------|
| <400> 163  |            |             |            |             |             |      |
| gccacctggc | cctcctgatc | gacgacacac  | gcacttgaaa | cttgtttctca | gggtgtgtgg  | 60   |
| aatcaacttt | ccggaagcaa | ccagcccacc  | agaggaggtc | ccgagcgcg   | gcggagacga  | 120  |
| tgcagcggag | actggttcag | cagtggagcg  | tcgcggtggt | cctgctgagc  | tacgcggtgc  | 180  |
| cctcctgcgg | gcgctcggtg | gaggggtctca | gccgccgcct | caaaagagct  | gtgtctgaac  | 240  |
| atcagctcct | ccatgacaag | gggaagtcca  | tccaagattt | acggcgacga  | ttcttccctc  | 300  |
| accatctgat | cgcagaaatc | cacacagctg  | aaatcagagc | tacctcgag   | gtgtccccta  | 360  |
| actccaagcc | ctctcccaac | acaaagaacc  | accccgctcg | atttggggtct | gatgatgagg  | 420  |
| gcagatacct | aactcaggaa | actaacaagg  | tggagacgta | caaagagcag  | ccgctcaaga  | 480  |
| cacctgggaa | gaaaaagaaa | ggcaagcccg  | ggaaacgcaa | ggagcaggaa  | aagaaaaaac  | 540  |
| ggcgaactcg | ctctgcctgg | ttagactctg  | gagtgactgg | gagtgaggta  | gaagggggacc | 600  |
| acctgtctga | cacctccaca | acgtcgctgg  | agctcgattc | acggaggcat  | tgaaattttc  | 660  |
| agcagagacc | ttccaaggac | atattgcagg  | attctgtaat | agtgaacata  | tggaaagtat  | 720  |
| tagaaatatt | tattgtctgt | aaatactgta  | aatgcattgg | aataaaaactg | tctcccccat  | 780  |
| tgctctatga | aactgcacat | tggtcattgt  | gaatatTTTT | TTTTTgcca   | aggctaatec  | 840  |
| aattattatt | atcacattta | ccataattta  | TTTTgtccat | tgatgtattt  | TTTTgtaaa   | 900  |
| tgtatcttgg | tgctgctgaa | TTTctatatt  | TTTTgtaaca | taattgcatt  | tagatataca  | 960  |
| tatcaagtat | gttgataaat | gacacaatga  | agtgctctca | TTTTgtggtt  | gatttttaatg | 1020 |
| aatgcctaaa | tataattatc | caaattgatt  | ttcctttgtg | catgtaaaaa  | taacagtatt  | 1080 |
| ttaaatttgt | aaagaatgtc | taataaaaata | taatctaatt | acatcatg    |             | 1128 |

<210> 164  
<211> 1310  
<212> DNA

<213> Homo sapiens

<400> 164

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gagacgtgta aacacactac ttatcattga tgcataatata aaaccatttt attttcgcta 180
ttattttcaga ggaagcgccct ctgattttgt tcttttttcc ctttttgctc tttctggcgtg 240
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gtctgaacat cagctcctcc atgacaaggg gaagtccatc caagatttac ggcgacgatt 480
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<210> 165

<211> 177

<212> PRT

<213> Homo sapiens

<400> 165

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          20          25          30
Arg Leu Lys Arg Ala Val Ser Glu His Gln Leu Leu His Asp Lys Gly
          35          40          45
Lys Ser Ile Gln Asp Leu Arg Arg Arg Phe Phe Leu His His Leu Ile
          50          55          60
Ala Glu Ile His Thr Ala Glu Ile Arg Ala Thr Ser Glu Val Ser Pro
65          70          75          80
Asn Ser Lys Pro Ser Pro Asn Thr Lys Asn His Pro Val Arg Phe Gly
          85          90          95
Ser Asp Asp Glu Gly Arg Tyr Leu Thr Gln Glu Thr Asn Lys Val Glu
          100         105         110
Thr Tyr Lys Glu Gln Pro Leu Lys Thr Pro Gly Lys Lys Lys Lys Gly
          115         120         125
Lys Pro Gly Lys Arg Lys Glu Gln Glu Lys Lys Lys Arg Arg Thr Arg
          130         135         140
Ser Ala Trp Leu Asp Ser Gly Val Thr Gly Ser Gly Leu Glu Gly Asp
145         150         155         160
His Leu Ser Asp Thr Ser Thr Thr Ser Leu Glu Leu Asp Ser Arg Arg
          165         170         175

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His

<210> 166  
 <211> 177  
 <212> PRT  
 <213> Homo sapiens

<400> 166  
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 Arg Leu Lys Arg Ala Val Ser Glu His Gln Leu Leu His Asp Lys Gly  
 35 40 45  
 Lys Ser Ile Gln Asp Leu Arg Arg Arg Phe Phe Leu His His Leu Ile  
 50 55 60  
 Ala Glu Ile His Thr Ala Glu Ile Arg Ala Thr Ser Glu Val Ser Pro  
 65 70 75 80  
 Asn Ser Lys Pro Ser Pro Asn Thr Lys Asn His Pro Val Arg Phe Gly  
 85 90 95  
 Ser Asp Asp Glu Gly Arg Tyr Leu Thr Gln Glu Thr Asn Lys Val Glu  
 100 105 110  
 Thr Tyr Lys Glu Gln Pro Leu Lys Thr Pro Gly Lys Lys Lys Gly  
 115 120 125  
 Lys Pro Gly Lys Arg Lys Glu Gln Glu Lys Lys Lys Arg Arg Thr Arg  
 130 135 140  
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 145 150 155 160  
 His Leu Ser Asp Thr Ser Thr Thr Ser Leu Glu Leu Asp Ser Arg Arg  
 165 170 175  
 His

<210> 167  
 <211> 3362  
 <212> DNA  
 <213> Homo sapiens

<400> 167  
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 tccatacacc ctacaataca gaggggtgtgg aaaagaggga aaatacattc atttcacacc 480  
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 ccatgaatgg gccacacctc gttgggggtgt gttcgatgag tataacaatg acaaaccttt 600  
 ctacataaat gggcaaaatc aaattaaagt gacaagggtgt tcatctgaca tcacaggcat 660  
 ttttgtgtgt gaaaaagggtc cttgccccca agaaaactgt attattagta agcttttttaa 720

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agaaggatgc acctttatct acaatagcac caaaaatgca actgcatcaa taatgttcat 780
gcaaagttaa tcttctgtgg ttgaattttg taatgcaagt acccacaacc aagaagcacc 840
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<210> 169

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<212> PRT

<213> Homo sapiens

<400> 169

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 180 185 190  
 Val Thr Arg Cys Ser Ser Asp Ile Thr Gly Ile Phe Val Cys Glu Lys  
 195 200 205  
 Gly Pro Cys Pro Gln Glu Asn Cys Ile Ile Ser Lys Leu Phe Lys Glu  
 210 215 220

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| Gly 225 | Cys | Thr | Phe | Ile     | Tyr 230 | Asn | Ser | Thr | Gln     | Asn 235 | Ala | Thr | Ala | Ser | Ile 240 |
| Met     | Phe | Met | Gln | Ser 245 | Leu     | Ser | Ser | Val | Val 250 | Glu     | Phe | Cys | Asn | Ala | Ser 255 |
| Thr     | His | Asn | Gln | Glu 260 | Ala     | Pro | Asn | Leu | Gln 265 | Asn     | Gln | Met | Cys | Ser | Leu 270 |
| Arg     | Ser | Ala | Trp | Asp 275 | Val     | Ile | Thr | Asp | Ser 280 | Ala     | Asp | Phe | His | His | Ser 285 |
| Phe     | Pro | Met | Asn | Gly 290 | Thr     | Glu | Leu | Pro | Pro 295 | Pro     | Pro | Thr | Phe | Ser | Leu 300 |
| Val 305 | Glu | Ala | Gly | Asp 310 | Lys     | Val | Val | Cys | Leu 315 | Val     | Leu | Asp | Val | Ser | Ser 320 |
| Lys     | Met | Ala | Glu | Ala 325 | Asp     | Arg | Leu | Leu | Gln 330 | Leu     | Gln | Gln | Ala | Ala | Glu 335 |
| Phe     | Tyr | Leu | Met | Gln 340 | Ile     | Val | Glu | Ile | His 345 | Thr     | Phe | Val | Gly | Ile | Ala 350 |
| Ser     | Phe | Asp | Ser | Lys 355 | Gly     | Glu | Ile | Arg | Ala 360 | Gln     | Leu | His | Gln | Ile | Asn 365 |
| Ser     | Asn | Asp | Asp | Arg 370 | Lys     | Leu | Leu | Val | Ser 375 | Tyr     | Leu | Pro | Thr | Thr | Val 380 |
| Ser 385 | Ala | Lys | Thr | Asp 390 | Ile     | Ser | Ile | Cys | Ser 395 | Gly     | Leu | Lys | Lys | Gly | Phe 400 |
| Glu     | Val | Val | Glu | Lys 405 | Leu     | Asn | Gly | Lys | Ala 410 | Tyr     | Gly | Ser | Val | Met | Ile 415 |
| Leu     | Val | Thr | Ser | Gly 420 | Asp     | Asp | Lys | Leu | Leu 425 | Gly     | Asn | Cys | Leu | Pro | Thr 430 |
| Val     | Leu | Ser | Ser | Gly 435 | Ser     | Thr | Ile | His | Ser 440 | Ile     | Ala | Leu | Gly | Ser | Ser 445 |
| Ala     | Ala | Pro | Asn | Leu 450 | Glu     | Glu | Leu | Ser | Arg 455 | Leu     | Thr | Gly | Gly | Leu | Lys 460 |
| Phe 465 | Phe | Val | Pro | Asp 470 | Ile     | Ser | Asn | Ser | Asn 475 | Ser     | Met | Ile | Asp | Ala | Phe 480 |
| Ser     | Arg | Ile | Ser | Ser 485 | Gly     | Thr | Gly | Asp | Ile 490 | Phe     | Gln | Gln | His | Ile | Gln 495 |
| Leu     | Glu | Ser | Thr | Gly 500 | Glu     | Asn | Val | Lys | Pro 505 | His     | His | Gln | Leu | Lys | Asn 510 |
| Thr     | Val | Thr | Val | Asp 515 | Asn     | Thr | Val | Gly | Asn 520 | Asp     | Thr | Met | Phe | Leu | Val 525 |
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| Thr     | Ala | Ser | Leu | Trp 565 | Ile     | Pro | Gly | Thr | Ala 570 | Lys     | Pro | Gly | His | Trp | Thr 575 |
| Tyr     | Thr | Leu | Asn | Asn 580 | Thr     | His | His | Ser | Leu 585 | Gln     | Ala | Leu | Lys | Val | Thr 590 |
| Val     | Thr | Ser | Arg | Ala 595 | Ser     | Asn | Ser | Ala | Val 600 | Pro     | Pro | Ala | Thr | Val | Glu 605 |
| Ala     | Phe | Val | Glu | Arg 610 | Asp     | Ser | Leu | His | Phe 615 | Pro     | His | Pro | Val | Met | Ile 620 |
| Tyr 625 | Ala | Asn | Val | Lys 630 | Gln     | Gly | Phe | Tyr | Pro 635 | Ile     | Leu | Asn | Ala | Thr | Val 640 |
| Thr     | Ala | Thr | Val | Glu 645 | Pro     | Glu | Thr | Gly | Asp 650 | Pro     | Val | Thr | Leu | Arg | Leu 655 |



Leu Asp Asp Gly Ala Gly Ala Asp Val Ile Lys Asn Asp Gly Ile Tyr  
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 Ser Arg Tyr Phe Phe Ser Phe Ala Ala Asn Gly Arg Tyr Ser Leu Lys  
                   675                  680                  685  
 Val His Val Asn His Ser Pro Ser Ile Ser Thr Pro Ala His Ser Ile  
                   690                  695                  700  
 Pro Gly Ser His Ala Met Tyr Val Pro Gly Tyr Thr Ala Asn Gly Asn  
 705                  710                  715                  720  
 Ile Gln Met Asn Ala Pro Arg Lys Ser Val Gly Arg Asn Glu Glu Glu  
                   725                  730                  735  
 Arg Lys Trp Gly Phe Ser Arg Val Ser Ser Gly Gly Ser Phe Ser Val  
                   740                  745                  750  
 Leu Gly Val Pro Ala Gly Pro His Pro Asp Val Phe Pro Pro Cys Lys  
                   755                  760                  765  
 Ile Ile Asp Leu Glu Ala Val Asn Arg Arg Gly Ile Asp Pro Ile Leu  
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<210> 171

<211> 1491

<212> DNA

<213> Homo sapiens

<400> 171

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<210> 172

<211> 364

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 172

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Leu Ala Ile Glu Ala Gly Phe His His Ile Asp Ser Ala His Val Tyr
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Gly Ser Val Lys Arg Glu Asp Ile Phe Tyr Thr Ser Lys Leu Trp Ser
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Asn Ser His Arg Pro Glu Leu Val Arg Pro Ala Leu Glu Arg Ser Leu
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Lys Asn Leu Gln Leu Asp Tyr Val Asp Leu Tyr Leu Ile His Phe Pro
145          150          155          160
Val Ser Val Lys Pro Gly Glu Glu Val Ile Pro Lys Asp Glu Asn Gly
          165          170          175
Lys Ile Leu Phe Asp Thr Val Asp Leu Cys Ala Thr Trp Glu Ala Met
          180          185          190
Glu Lys Cys Lys Asp Ala Gly Leu Ala Lys Ser Ile Gly Val Ser Asn
          195          200          205
Phe Asn His Arg Leu Leu Glu Met Ile Leu Asn Lys Pro Gly Leu Lys
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Tyr Lys Pro Val Cys Asn Gln Val Glu Cys His Pro Tyr Phe Asn Gln
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Tyr Ser Ala Leu Gly Ser His Arg Glu Glu Pro Trp Val Asp Pro Asn
          260          265          270
Ser Pro Val Leu Leu Glu Asp Pro Val Leu Cys Ala Leu Ala Lys Lys
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His Lys Arg Thr Pro Ala Leu Ile Ala Leu Arg Tyr Gln Leu Gln Arg
          290          295          300
Gly Val Val Val Leu Ala Lys Ser Tyr Asn Glu Gln Arg Ile Arg Gln
305          310          315          320
Asn Val Gln Val Phe Glu Phe Gln Leu Thr Ser Glu Glu Met Lys Ala
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Ile Asp Gly Leu Asn Arg Asn Val Arg Tyr Leu Thr Leu Asp Ile Phe
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&lt;210&gt; 173

&lt;211&gt; 1988

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 173

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&lt;210&gt; 174

&lt;211&gt; 238

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 174

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Gly Ala Ala Ser Pro Arg Pro Leu Arg Phe Cys Gly Gly Ala Arg Ala
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Arg Arg Pro Leu Ser Ala Val Ala Arg Pro Ala Arg Ser Ser Asp Pro
          20          25          30
Leu Arg Ser Ala Pro Leu Gly Pro Ala Pro Pro Val Asn Met Ile Arg
          35          40          45
Cys Gly Leu Ala Cys Glu Arg Cys Arg Trp Ile Leu Pro Leu Leu Leu
          50          55          60
Leu Ser Ala Ile Ala Phe Asp Ile Ile Ala Leu Ala Gly Arg Gly Trp

```

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| 65  |     | 70  |     | 75  |     | 80  |     |     |     |     |     |     |     |     |     |
| Leu | Gln | Ser | Ser | Asp | His | Gly | Gln | Thr | Ser | Ser | Leu | Trp | Trp | Lys | Cys |
|     |     |     |     | 85  |     |     |     |     | 90  |     |     |     |     | 95  |     |
| Ser | Gln | Glu | Gly | Gly | Gly | Ser | Gly | Ser | Tyr | Glu | Glu | Gly | Cys | Gln | Ser |
|     |     |     | 100 |     |     |     |     | 105 |     |     |     |     | 110 |     |     |
| Leu | Met | Glu | Tyr | Ala | Trp | Gly | Arg | Ala | Ala | Ala | Ala | Met | Leu | Phe | Cys |
|     |     |     | 115 |     |     |     | 120 |     |     |     |     | 125 |     |     |     |
| Gly | Phe | Ile | Ile | Leu | Val | Ile | Cys | Phe | Ile | Leu | Ser | Phe | Phe | Ala | Leu |
|     |     |     | 130 |     |     |     | 135 |     |     |     | 140 |     |     |     |     |
| Cys | Gly | Pro | Gln | Met | Leu | Val | Phe | Leu | Arg | Val | Ile | Gly | Gly | Leu | Leu |
| 145 |     |     |     |     | 150 |     |     |     |     | 155 |     |     |     |     | 160 |
| Ala | Leu | Ala | Ala | Val | Phe | Gln | Ile | Ile | Ser | Leu | Val | Ile | Tyr | Pro | Val |
|     |     |     |     | 165 |     |     |     |     | 170 |     |     |     |     | 175 |     |
| Lys | Tyr | Thr | Gln | Thr | Phe | Thr | Leu | His | Ala | Asn | Pro | Ala | Val | Thr | Tyr |
|     |     |     | 180 |     |     |     |     | 185 |     |     |     |     | 190 |     |     |
| Ile | Tyr | Asn | Trp | Ala | Tyr | Gly | Phe | Gly | Trp | Ala | Ala | Thr | Ile | Ile | Leu |
|     |     |     | 195 |     |     |     | 200 |     |     |     |     | 205 |     |     |     |
| Ile | Gly | Cys | Ala | Phe | Phe | Phe | Cys | Cys | Leu | Pro | Asn | Tyr | Glu | Asp | Asp |
|     |     |     | 210 |     |     |     | 215 |     |     |     | 220 |     |     |     |     |
| Leu | Leu | Gly | Asn | Ala | Lys | Pro | Arg | Tyr | Phe | Tyr | Thr | Ser | Ala |     |     |
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<210> 175  
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 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
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 4036, 4056, 4062, 4080, 4088, 4115  
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 ttactgtgtt tgtgtathtt aaaggcgaga agacgagggg aacaaaacca gctggatcca 180  
 tccatcaccg tgggtggttt taatttttctg ttttttctcg ttattttttt ttaaacaacc 240  
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&lt;210&gt; 176

&lt;211&gt; 579

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 176

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Met | Asn | Lys | Leu | Tyr | Ile | Gly | Asn | Leu | Ser | Glu | Asn | Ala | Ala | Pro | Ser |
| 1   |     |     |     | 5   |     |     |     |     | 10  |     |     |     |     | 15  |     |
| Asp | Leu | Glu | Ser | Ile | Phe | Lys | Asp | Ala | Lys | Ile | Pro | Val | Ser | Gly | Pro |
|     |     |     | 20  |     |     |     |     | 25  |     |     |     |     | 30  |     |     |
| Phe | Leu | Val | Lys | Thr | Gly | Tyr | Ala | Phe | Val | Asp | Cys | Pro | Asp | Glu | Ser |
|     |     | 35  |     |     |     |     | 40  |     |     |     |     | 45  |     |     |     |
| Trp | Ala | Leu | Lys | Ala | Ile | Glu | Ala | Leu | Ser | Gly | Lys | Ile | Glu | Leu | His |
|     | 50  |     |     |     |     | 55  |     |     |     |     | 60  |     |     |     |     |
| Gly | Lys | Pro | Ile | Glu | Val | Glu | His | Ser | Val | Pro | Lys | Arg | Gln | Arg | Ile |
| 65  |     |     |     | 70  |     |     |     |     |     | 75  |     |     |     | 80  |     |
| Arg | Lys | Leu | Gln | Ile | Arg | Asn | Ile | Pro | Pro | His | Leu | Gln | Trp | Glu | Val |
|     |     |     | 85  |     |     |     |     |     | 90  |     |     |     |     | 95  |     |
| Leu | Asp | Ser | Leu | Leu | Val | Gln | Tyr | Gly | Val | Val | Glu | Ser | Cys | Glu | Gln |
|     |     |     | 100 |     |     |     |     | 105 |     |     |     |     | 110 |     |     |
| Val | Asn | Thr | Asp | Ser | Glu | Thr | Ala | Val | Val | Asn | Val | Thr | Tyr | Ser | Ser |
|     |     | 115 |     |     |     |     | 120 |     |     |     |     | 125 |     |     |     |
| Lys | Asp | Gln | Ala | Arg | Gln | Ala | Leu | Asp | Lys | Leu | Asn | Gly | Phe | Gln | Leu |
|     | 130 |     |     |     |     | 135 |     |     |     |     | 140 |     |     |     |     |
| Glu | Asn | Phe | Thr | Leu | Lys | Val | Ala | Tyr | Ile | Pro | Asp | Glu | Met | Ala | Ala |
| 145 |     |     |     |     | 150 |     |     |     |     | 155 |     |     |     | 160 |     |
| Gln | Gln | Asn | Pro | Leu | Gln | Gln | Pro | Arg | Gly | Arg | Arg | Gly | Leu | Gly | Gln |
|     |     |     |     | 165 |     |     |     |     | 170 |     |     |     |     | 175 |     |
| Arg | Gly | Ser | Ser | Arg | Gln | Gly | Ser | Pro | Gly | Ser | Val | Ser | Lys | Gln | Lys |
|     |     | 180 |     |     |     |     |     | 185 |     |     |     |     | 190 |     |     |
| Pro | Cys | Asp | Leu | Pro | Leu | Arg | Leu | Val | Pro | Thr | Gln | Phe | Val | Gly |     |
|     |     | 195 |     |     |     |     | 200 |     |     |     | 205 |     |     |     |     |
| Ala | Ile | Ile | Gly | Lys | Glu | Gly | Ala | Thr | Ile | Arg | Asn | Ile | Thr | Lys | Gln |
|     | 210 |     |     |     |     | 215 |     |     |     |     | 220 |     |     |     |     |
| Thr | Gln | Ser | Lys | Ile | Asp | Val | His | Arg | Lys | Glu | Asn | Ala | Gly | Ala | Ala |
| 225 |     |     |     | 230 |     |     |     |     |     | 235 |     |     |     | 240 |     |
| Glu | Lys | Ser | Ile | Thr | Ile | Leu | Ser | Thr | Pro | Glu | Gly | Thr | Ser | Ala | Ala |
|     |     |     | 245 |     |     |     |     |     | 250 |     |     |     |     | 255 |     |
| Cys | Lys | Ser | Ile | Leu | Glu | Ile | Met | His | Lys | Glu | Ala | Gln | Asp | Ile | Lys |
|     |     | 260 |     |     |     |     | 265 |     |     |     |     | 270 |     |     |     |
| Phe | Thr | Glu | Glu | Ile | Pro | Leu | Lys | Ile | Leu | Ala | His | Asn | Asn | Phe | Val |
|     | 275 |     |     |     |     |     | 280 |     |     |     |     | 285 |     |     |     |
| Gly | Arg | Leu | Ile | Gly | Lys | Glu | Gly | Arg | Asn | Leu | Lys | Lys | Ile | Glu | Gln |
|     | 290 |     |     |     |     | 295 |     |     |     | 300 |     |     |     |     |     |
| Asp | Thr | Asp | Thr | Lys | Ile | Thr | Ile | Ser | Pro | Leu | Gln | Glu | Leu | Thr | Leu |
| 305 |     |     |     | 310 |     |     |     |     |     | 315 |     |     |     | 320 |     |
| Tyr | Asn | Pro | Glu | Arg | Thr | Ile | Thr | Val | Lys | Gly | Asn | Val | Glu | Thr | Cys |
|     |     |     | 325 |     |     |     |     |     | 330 |     |     |     |     | 335 |     |
| Ala | Lys | Ala | Glu | Glu | Glu | Ile | Met | Lys | Lys | Ile | Arg | Glu | Ser | Tyr | Glu |
|     |     | 340 |     |     |     |     | 345 |     |     |     |     | 350 |     |     |     |
| Asn | Asp | Ile | Ala | Ser | Met | Asn | Leu | Gln | Ala | His | Leu | Ile | Pro | Gly | Leu |
|     |     | 355 |     |     |     |     | 360 |     |     |     |     | 365 |     |     |     |
| Asn | Leu | Asn | Ala | Leu | Gly | Leu | Phe | Pro | Pro | Thr | Ser | Gly | Met | Pro | Pro |
|     | 370 |     |     |     |     | 375 |     |     |     |     | 380 |     |     |     |     |
| Pro | Thr | Ser | Gly | Pro | Pro | Ser | Ala | Met | Thr | Pro | Pro | Tyr | Pro | Gln | Phe |
| 385 |     |     |     | 390 |     |     |     |     |     | 395 |     |     |     | 400 |     |

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<210> 177
<211> 401
<212> DNA
<213> Homo sapiens
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<210> 178
<211> 561
<212> DNA
<213> Homo sapiens
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| acgcctttca  | agggtgtacg  | caaagcactc | attgataccc | ttttggatgg | ctatgaaaca | 60  |
| gcccgcctatg | ggacaggggt  | ctttggccag | aatgagtacc | tacgctatca | ggaggccctg | 120 |
| agtgagctgg  | ccactgcggt  | taaagcacga | attgggagct | ctcagcgaca | tcaccagtca | 180 |
| gcagccaaaag | acctaactca  | gtcccctgag | gtctcccaa  | caaccatcca | ggtgacatac | 240 |
| ctcccctcca  | gtcagaagag  | taaacgtgcc | aagcacttcc | ttgaattgaa | gagctttaag | 300 |
| gataactata  | acacattgga  | gagtactctg | tgacggagct | gaaggactct | tgccgtagat | 360 |
| taagccagtc  | agttgcaatg  | tgcaagacag | gctgcttgcc | gggccgccct | cggaacatct | 420 |
| ggcccagcag  | gcccaagactg | tatccatcca | agttcccgtt | gtatccagag | ttcttagagc | 480 |

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ttgtgtctaa agggtaatc cccaaccctt ccttatgagc atttttagaa cattggctaa 540
gactattttc cccagtagc g                                     561

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<210> 179

<211> 521

<212> DNA

<213> Homo sapiens

<400> 179

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cccaacgcgt ttgcaaata tcccctggta gcctacttcc ttacccccga atattggtaa 60
gatcgagcaa tggcttcagg acatgggttc tcttctcctg tgatcattca agtgctcact 120
gcatgaagac tggcttgtct cagtgtttca acctcaccag ggctgtctct tgggccacac 180
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aggataagtg ggatctacca attgattctg gcaaaacaat ttctaagatt tttttgcttt 480
atgtgggaaa cagatctaaa tctcatttta tgctgtattt t                                     521

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<210> 180

<211> 417

<212> DNA

<213> Homo sapiens

<400> 180

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tcctgggccg cctggcgccg atcgtggcta aacagggtact gctgggcccg aagggtggtg 120
tcgtacgctg tgaaggcatc aacatttctg gcaatttcta cagaaacaag ttgaagtacc 180
tggcttttct ccgcaagcgg atgaacacca accttcccg agggccctac cacttccggg 240
ccccagccg catcttctgg cggaccgtgc gaggtatgct gccccacaaa accaagcgag 300
gccaggccgc tctggaccgt ctcaagggtg ttgacggcat cccaccgccc tacgacaaga 360
aaaagcggat ggtggttctt gctgccctca aggtcgtgcg tctgaagcct acaagaa 417

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<210> 181

<211> 283

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 35

<223> n = A,T,C or G

<400> 181

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caagaactca agtgtaactg tgataaaata acctttccca ggtatattgg cagggtatgtg 120
tgtaatctca gaatacacag gtgacataga tatgatatga caactggtaa tgggtggattc 180
atttacattg ttacacttc tatgaccagg ccttaaggga aggtcagttt tttaaaaaac 240
caagtagtgt ctctctacct atctccagat acatgtcaaa aaa                                     283

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<210> 182

<211> 401

<212> DNA

<213> Homo sapiens



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<400> 182
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tatttcccac agtgaaagaa aacgctggcc tatcagttac attacaaaag gcagatttca 120
agaggattga gtaagtagtt ggatggcttt cataaaaaaca agaattcaag aagaggattc 180
atgctttaag aaacatttgt tatacattcc tcacaaatta tacctgggat aaaactatg 240
tagcaggcag tgtgttttcc ttccatgtct ctctgcacta cctgcagtgt gtcctctgag 300
gctgcaagtc tgtcctatct gaattcccag cagaagcact aagaagctcc accctatcac 360
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<210> 183

<211> 366

<212> DNA

<213> Homo sapiens

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<221> misc\_feature

<222> 325

<223> n = A,T,C or G

<400> 183

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tttaaggaca aagatgaagt cactgtaaac taatctgtca ttgtttttac ctcccttttc 180
tttttcagtg cagaaattaa aagtaagtat aaagcacctg gattgggagt gtttttgcgt 240
gtgtcggaat cactggtaaa tgttggctga gaacaatccc tccccttgca cttgtgaaaa 300
cactttgagc gctttaagag attancctga gaaataatta aatatctttt ctcttcaaaa 360
aaaaaa 366

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<210> 184

<211> 370

<212> DNA

<213> Homo sapiens

<400> 184

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taaaatgtta gtctacatag atgggtgatt gtaactttat tgccattaaa agatttcaaa 180
ttgcattcat gcttctgtgt acacataatg aaaaatgggc aaataatgaa gatctctcct 240
tcagtctgct ctgtttaatt ctgctgtctg ctcttctcta atgctgcgtc cctaattgta 300
cacagtttag tgatatctag gagtataaag ttgtcgccca tcaataaaaa tcacaaagtt 360
ggtttaaaaa 370

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<210> 185

<211> 107

<212> DNA

<213> Homo sapiens

<400> 185

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ctcatattat tttccttttg agaaattgga aactctttct gttgctatta tattaataaa 60
gttggtgttt attttctggg agtcaccttc cccatttaaa aaaaaaa 107

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<210> 186

<211> 309

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<212> DNA

<213> Homo sapiens

<400> 186

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gaaaggatgg ctctggttgc cacagagctg ggacttcacg ttcttctaga gagggccaca 60
agagggccac aggggtggcc gggagtgtgc agctgatgcc tgctgagagg caggaattgt 120
gccagtgagt gacagtcacg agggagtgtc tcttcttggg gaggaagaa ggtagagcct 180
ttctgtctga atgaaaggcc aaggctacag tacaggggcc cgcccagcc aggtgtgtta 240
tgcccacgta gtggaggcct ctggcagatc ctgcattcca aggtcactgg actgtacgtt 300
tttatggtt                                     309

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<210> 187

<211> 477

<212> DNA

<213> Homo sapiens

<400> 187

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ttcagtccta gcaagaagcg agaattctga gatcctccag aaagtcgagc agcaccacc 60
tccaacctcg ggccagtgtc ttcaggtctt actggggacc tgcgagctgg cctaattgtg 120
tggcctgcaa gccaggccat ccctgggccc cacagacgag ctccgagcca ggtcaggctt 180
cggaggccac aagctcagcc tcaggcccag gcactgattg tggcagaggg gccactacc 240
aaggtctagc taggcccag acctagttac ccagacagtg agaagcccct ggaaggcaga 300
aaagtgtgga gcatggcaga cagggaaggg aaacattttc agggaaaaga catgtatcac 360
atgtcttcag aagcaagtca ggtttcatgt aaccgagtgt cctcttgctg gtccaaaagt 420
agcccagggc tgtagcacag gcttcacagt gattttgtgt tcagccgtga gtcacac 477

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<210> 188

<211> 220

<212> DNA

<213> Homo sapiens

<400> 188

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taaatatggt agatattaat attcctctta gatgaccagt gattccaatt gtcccaagtt 60
ttaaataagt accctgtgag tatgagataa attagtgcac atcagaacaa gtttcagtat 120
cagatgttca agaggaagtt gctattgcat tgattttaat atttgtacat aaacactgat 180
ttttttgagc attattttgt atttgttgta ctttaataacc                220

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<210> 189

<211> 417

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 76, 77

<223> n = A,T,C or G

<400> 189

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accatcttga cagaggatag atgctcccaa aacgtttgtt accacactta aaaatcactg 60
ccatcattaa gcatcnnttt caaaattata gccattcatg atttactttt tccagatgac 120
tatcattatt ctagtctttt gaatttgtaa ggggaaaaaa aacaaaaaca aaaacttacg 180
atgcactttt ctccagcaca tcagatttca aattgaaaat taaagacatg ctatggtaat 240
gcacttgcta gtactacaca ctttgtacaa caaaaaacag aggcaagaaa caacggaaag 300
agaaaagcct tcctttgttg gcccttaaac tgagtcaaga tctgaaatgt agagatgac 360

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tctgacgata cctgtatgtt cttatttgtt aaataaaatt gctggatatga aatgaca 417

<210> 190

<211> 497

<212> DNA

<213> Homo sapiens

<400> 190

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gcaactgcggc gctctcccgt cccgcgggtg ttgctgctgc tgccgctgct gctgggcctg 60
aacgcaggag ctgtcattga ctggcccaca gaggagggca aggaagtatg ggattatgtg 120
acggtcgcga aggatgccta catgttcttg tggctctatt atgccaccaa ctccctgcaag 180
aactttctcag aactgcccct ggtcatgtgg cttcagggcg gtccaggcgg ttctagcact 240
ggatttggaa actttgagga aattggggcc cttgacagtg atctcaaacc acggaaaacc 300
acctggctcc aggcgtgccag tctcctatct gtggataatc ccgtgggcac tgggttcagt 360
tatgtgaatg gtagtgtgct ctatgccaa gacctggcta tgggtggctc agacatgatg 420
gttctcctga agaccttctt cagttgccac aaagaattcc agacagttcc attctacatt 480
ttctcagagt cctatgg                                     497

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<210> 191

<211> 175

<212> DNA

<213> Homo sapiens

<400> 191

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atgttgaata ttttgcttat taactttgtt tattgtcttc tccctcgatt agaataattag 60
ctacttgagt acaaggattt gagcctgtta cattcactgc tgaatttttag gtcctcggaa 120
gatacccagc attcaataga gaccacacaa taaatatatg tcaaataaaa aaaaaa 175

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<210> 192

<211> 526

<212> DNA

<213> Homo sapiens

<400> 192

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agtaaacatt attatTTTTT ttatatTTTgc aaaggaaaca tatctaatec ttccatataga 60
aagaacagta ttgctgtaat tctTTTTctt ttcttctca tttcctctgc cccttaaaag 120
attgaagaaa gagaaacttg tcaactcata tccacgttat ctagcaaagt acataagaat 180
ctatcactaa gtaatgtatc cttcagaatg tgttggttta ccagtgcacac cccatattca 240
tcacaaaatt aaagcaagaa gtccatagta atttatttgc taatagtggg tttttaatgc 300
tcagagtttc tgagggtcaaa ttttatcttt tcacttaca gctctatgat cttaaataat 360
ttacttaatg tatTTTgggtg tatTTTcttc aaattaatat tgggtgttcaa gactatatct 420
aatcctctg atcactttga gaaacaaact tttattaaat gtaaggcact tttctatgaa 480
ttttaaatat aaaaataaat attgttctga ttattactga aaaaaa 526

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<210> 193

<211> 553

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 290, 300, 411, 441

<223> n = A,T,C or G

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<400> 193
tccattgtgg tggaaattcgc tctctggtaa aggcgtgcag gtgttggccg cggcctctga 60
gctgggatga gccgtgctcc cgggtggaagc aagggagccc agccggagcc atggccagta 120
cagtggtagc agttggactg accattgctg ctgcaggatt tgcaggccgt tacgttttgc 180
aagccatgaa gcataatggag cctcaagtaa aacaagtttt tcaaagccta ccaaaatctg 240
ccttcagtgg tggctattat agaggtgggt ttgaacccaa aatgacaaan cgggaagcan 300
cattaatact aggtgtaagc cctactgcc aataaaggaa aataagagat gctcatcgac 360
gaattatgct tttaaatcat cctgacaaag gaggatctcc ttatatagca nccaaaatca 420
atgaagctaa agatttacta naaggatcaag ctaaaaaatg aagtaaatgt atgatgaatt 480
ttaagttcgt attagtttat gtatatgagt actaagtttt tataataaaa tgcctcagag 540
ctacaatttt aaa 553

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<210> 194
<211> 320
<212> DNA
<213> Homo sapiens

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<400> 194
cccttcccaa tccatcagta aagaccccat ctgccttgtc catgccgttt cccaacaggg 60
atgtcacttg atatgagaat ctcaaatctc aatgccttat aagcattcct tctgtgtcc 120
attaagactc tgataattgt ctccctcca taggaatttc tcccaggaaa gaaatatatc 180
cccatctccg ttctatatca gaactaccgt ccccgatatt cccttcagag agattaaaga 240
ccagaaaaaa gtgagcctct tcatctgcac ctgtaatagt ttcagttcct attttcttcc 300
attgacccat atttatacct 320

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<210> 195
<211> 320
<212> DNA
<213> Homo sapiens

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<220>
<221> misc_feature
<222> 203, 218
<223> n = A,T,C or G

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<400> 195
aagcatgacc tggggaaatg gtcagacctt gtatttgtgt tttggccttg aaagtagcaa 60
gtgaccagaa tctgccatgg caacaggctt taaaaaagac ccttaaaaag acactgtctc 120
aactgtggtg ttagcaccag ccagctctct gtacatttgc tagcttgtag ttttctaaga 180
ctgagtaaac ttcttatttt tanaaaagggg aggcctggntt gtaactttcc ttgtacttaa 240
ttgggtaaaa gtcttttcca caaaccacca tctattttgt gaactttgtt agtcatcttt 300
tatttggtaa attatgaact 320

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<210> 196
<211> 357
<212> DNA
<213> Homo sapiens

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<220>
<221> misc_feature
<222> 36
<223> n = A,T,C or G

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<400> 196

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atataaaata atacgaaact ttaaaaagca ttggantgtc agtatgttga atcagtagtt 60
tcactttaac tgtaaacaat ttcttaggac accatttggg ctagtttctg tgtaagtgtg 120
aatactacaa aaacttattt atactgttct tatgtcattt gttatattca tagatttata 180
tgatgatatg acatctggct aaaaagaaat tattgcaaaa ctaaccacta tgtacttttt 240
tataaatact gtatggacaa aaaatggcat tttttatatt aaattgttta gctctggcaa 300
aaaaaaaaa ttttaagagc tgggtactaat aaaggattat tatgactgtt aaaaaaa 357

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<210> 197

<211> 565

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 27

<223> n = A,T,C or G

<400> 197

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tcagctgagt accatcagga tatttanccc ttttaagtgtc gttttgggag tagaaaacta 60
aagcaacaat acttcctctt gacagctttg attggaatgg ggttattaga tcattcacct 120
tggtcctaca ctttttagga tgcttggtga acataacacc acttataatg aacatccctg 180
gttcctatat tttgggctat gtgggtagga attgttactt gttactgcag cagcagccct 240
agaaagtaag cccagggctt cagatctaag ttagtccaaa agctaaatga tttaaagtca 300
agttgtaatg ctaggcataa gcactctata atacattaaa ttataggccg agcaattagg 360
gaatgtttct gaaacattaa acttgtatct atgtcactaa aattctaaca caaacttaaa 420
aaatgtgtct catacatatg ctgtactagg ctccatcatg catttctaaa tttgtgtatg 480
atattgaatat atgaaagaat ttatacaaga gtgttattta aaattattaa aaataaatgt 540
atataatttg tacctattgt aaaaaa

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<210> 198

<211> 484

<212> DNA

<213> Homo sapiens

<400> 198

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tatgtaagta ttggtgtctg ctttaaaaaa ggagaccag acttcacctg tccttttttaa 60
acatttgaga acagtgttac tctgagcagt tgggccacct tcaccttatc cgacagctga 120
ctggttgatg tgtccattgt cgccagtttg gctgttgccc ggacaggaca ggacctccat 180
tgggcgcagc agcagggtggc aggggtgtgg cttgaggtgg gtggcagcgt ctggtcctcc 240
tctctggtgc tttctgagag ggtctctaaa gcagagtgtg gttggcctgg gggaaggcag 300
agcacgtatt tctccccctc agtacctctg catttgtagg tgttccctct ggctttctga 360
agggcagcag actcttgagt atactgcaga ggacatgctt tatcagtagg tcctgagggc 420
tccaggggct caactgacca agtaacacag aagttggggg atgtggccta tttgggtcgg 480
aaac

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<210> 199

<211> 429

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 77, 88, 134, 189, 227, 274, 319

<223> n = A,T,C or G

&lt;400&gt; 199

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gcttatgttt tttgttttaa cttttgtttt ttaacattta gaatattaca ttttgtatta 60
tacagtacct ttctcanaca ttttgtanaa ttcatttcgg cagctcacta ggattttgct 120
gaacattaaa aagngtgata gcgatattag ngccaatcaa atggaaaaaa ggtagtctta 180
ataaacaana cacaacgttt ttatacaaca tactttaaaa tattaanaaa actccttaat 240
attgtttcct attaagtatt attccttggg caanattttc tgatgctttt gattttctct 300
caatttagca tttgctttng gtttttttct ctatttagca ttctgttaag gcacaaaaac 360
tatgtactgt atgggaaatg ttgtaaatat taccttttcc acatttttaa cagacaactt 420
tgaatccaa                                     429

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&lt;210&gt; 200

&lt;211&gt; 279

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 200

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gcttttttga ggaattacag ggaagctcct ggaattgtac atggatatct ttatccctag 60
ggggaaatca aggagctggg caccctaat tctttatgga agtggtttaa actattttaa 120
ttttattaca agtattacta gagtagtggt tctactctaa gatttcaaaa gtgcatttaa 180
aatcatacat gttcccgctt gcaaatatat tgttattttg gtggagaaaa aaatagtata 240
ttctacataa aaaattaaag atattaacta agaaaaaaaa 279

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&lt;210&gt; 201

&lt;211&gt; 569

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 201

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taggtcagta tttttagaaa ctcttaatag ctcatactct tgataccaaa agcagccctg 60
attgttaaag cacacacctg cacaagaagc agtgatgggt gcatttacat ttccctgggtg 120
cacaaaaaaa aattctcaaa aagcaaggac ttacgctttt tgcaaagcct ttgagaagtt 180
actggatcat aggaagctta taacaagaat ggaagattct taaataactc actttctttg 240
gtatccagta acagtagatg ttcaaaaatat gtagctgatt aataccagca ttgtgaacgc 300
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aattaatgtt atttatacac tgcccttccat gacttttact ttgccctaag ctaatctcca 420
aaatctgaaa tgctactcca atatcagaaa aaaaggggga ggtggaatta tatttctctgt 480
gatttttaaga gtacagagaa tcatgcacat ctctgattag ttcatatatg tctagtgtgt 540
aataaaaagtc aaagatgaac tctcaaaaa 569

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&lt;210&gt; 202

&lt;211&gt; 501

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 202

```

attaataggc ttaataattg ttggcaagga tccttttgct ttctttggca tgcaagctcc 60
tagcatctgg cagtggggcc aagaaaataa ggtttatgca tgtatgatgg ttttcttctt 120
gagcaacatg attgagaacc agtgatgtc aacagggtgca tttgagataa ctttaaatga 180
tgtacctgtg tgggtctaagc tggaatctgg tcaccttcca tccatgcaac aacttgttca 240
aattcttgac aatgaaatga agctcaatgt gcatatggat tcaatcccac accatcgatc 300
atagcaccac ctatcagcac tgaaaactct tttgcattaa gggatcattg caagagcagc 360
gtgactgaca ttatgaaggc ctgtactgaa gacagcaagc tgtagtagca gaccagatgc 420
tttcttggca ggctcgttgt acctcttggg aaacctcaat gcaagatagt gtttcagtgc 480

```

tgccatattt tggaattctg c

501

<210> 203

<211> 261

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 36, 96

<223> n = A,T,C or G

<400> 203

gacaagctcc tggctctgag atgtcttctc gttaangaga tgggcctttt ggaggtaaag 60  
gataaaatga atgagttctg tcatgattca ctattntata acttgcata cctttactgt 120  
gttagctctt tgaatgttct tgaattttta gactttcttt gtaaataaat gatatgtcct 180  
tatcattgta taaaagctgt tatgtgcaac agtgtggaga ttccttgtct gatttaataa 240  
aataacttaaa cactgaaaaa a 261

<210> 204

<211> 421

<212> DNA

<213> Homo sapiens

<400> 204

agcatctttt ctacaacgtt aaaattgcag aagtagctta tcattaaaaa acaacaacaa 60  
caacaataac aataaatcct aagtgtaaat cagttattct accccctacc aaggatatca 120  
gcctgttttt tccctttttt ctcttgggaa taattgtggg cttcttccca aatttctaca 180  
gcctctttcc tcttctcatg cttgagcttc cctgtttgca cgcattgcgtg tgcaggactg 240  
gcttgtgtgc ttggactcgg ctccaggtgg aagcatgctt tcccttgta ctgttggaga 300  
aactcaaac ttcaagccct aggtgtagcc attttgtcaa gtcatacaact gtatttttgt 360  
actggcatta acaaaaaaag aagataaaat attgtaccat taaactttta taaaacttta 420  
a 421

<210> 205

<211> 460

<212> DNA

<213> Homo sapiens

<400> 205

tactctcaca atgaaggacc tggaaatgaaa aatctgtgtc taaacaagtc ctcttttagat 60  
tttagtgcaa atccagagacc agcgtcgggt gcctcgagta attctttcat gggtagcttt 120  
ggaaaagctc tcaggagacc tcacctagat gcctattcaa gctttggaca gccatcagat 180  
tgtcagccaa gagcctttta ttgaaaagct cattcttccc cagacttggg ctctgggtca 240  
gaggaagatg ggaaagaaa gacagatttt caggaagaaa atcacatttg tacctttaaa 300  
cagacttttag aaaactacag gactccaaat tttcagtcct atgacttggg cacatagact 360  
gaatgagacc aaaggaaaag cttaacatac tacctcaagg tgaactttta tttaaaagag 420  
agagaatctt atgtttttta aatggagtta tgaattttta 460

<210> 206

<211> 481

<212> DNA

<213> Homo sapiens

&lt;400&gt; 206

```

tgtggtggaa ttcgggacgc cccagaccc tgactttttc ctgctggtggc cgtctcctcc 60
tgcggaagca gtgacctctg acccctggtg accttcgctt tgagtgcctt ttgaacgctg 120
gtcccgcggg acttggtttt ctcaagctct gtctgtccaa agacgctccg gtcgagggtcc 180
cgctgtccct ggggtggatac ttgaacccca gacgcccctc tgtgctgctg tgtccggagg 240
cggccttccc atctgcctgc ccacccggag ctctttccgc cggcgcaggg tcccaagccc 300
acctcccgcc ctacgtcctg cgggtgtgct ctgggcacgt cctgcacaca caatgcaagt 360
cctggcctcc gcgcccggcc gccacgcga gccgtaccgc ccgccaactc tgttatttat 420
ggtgtgaccc cctggagggt ccctcgcccc accggggcta ttattgttt aatttatttg 480
t 481

```

&lt;210&gt; 207

&lt;211&gt; 605

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 207

```

accttttttg gattcagggc tctcacaat taaaatgagt gtaatgaaac aagggtgaaaa 60
tatagaagca tccctttgta tactgttttg ctacttacag tgtacttggc attgctttat 120
ctcactggat tctcacggta ggatttctga gatcttaatc taagctccaa agttgtctac 180
ttttttgatc ctagggtgct ccttttgttt tacagagcag ggtcacttga tttgctagct 240
gggtggcagaa ttggcaccat taccaggtc tgactgacca ccagtcagag gcactttatt 300
tgtatcatga aatgatttga aatcatttga aagcagcgaa gtctgataat gaatgccagc 360
tttccttggtg ctttgataac aaagactcca aatattctgg agaacctgga taaaagtttg 420
aagggtctaga ttgggatttg aagacaaaat tgtaggaaat cttacatttt tgcaataaca 480
aacattaatg aaagcaaaac attataaaag taattttaat tcaccacata cttatcaatt 540
tcttgatgct tccaaatgac atctaccaga tatggttttg tggacatctt tttctgttta 600
cataa 605

```

&lt;210&gt; 208

&lt;211&gt; 655

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 208

```

ggcgttggtt tggattcccc tcgtaactta aagggaaact ttcacaatgt ccggagccct 60
tgatgtcctg caaatgaagg aggaggatgt ccttaagttc cttgcagcag gaaccactt 120
aggtggcacc aatcttgact tccagatgga acagtacatc tataaaagga aaagtgatgg 180
catctatatc ataaatctca agaggacctg ggagaagctt ctgctggcag ctgctgcaat 240
tgttgccatt gaaaaccctg ctgatgtcag tgttatatcc tccaggaata ctggccagag 300
ggctgtgctg aagtttgctg ctgccactgg agccactcca attgctggcc gcttacttcc 360
tggaaccttc actaaccaga tccaggcagc ctccgggag ccacggcttc ttgtggttac 420
tgaccccagg gctgaccacc agcctctcac ggaggcatct tatgttaacc tacctacct 480
tgcgctgtgt aacacagatt ctccctctgc ctatgtggac attgccatcc catgcaaca 540
caaggagct cactcagtg gtttgatgt gtggatgctg gtcgggaag tctgcgcat 600
gcgtggcacc atttcccggtg aacacccatg ggaggcatg cctgatctgt acttc 655

```

&lt;210&gt; 209

&lt;211&gt; 621

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 209

```

catttagaac atggttatca tccaagacta ctctaccctg caacattgaa ctcccaagag 60

```



```

caaatccaca ttcctcttga gttctgcagc ttctgtgtaa atagggcagc tgcgtctat 120
gccgtagaat cacatgatct gaggaccatt catggaagct gctaaatagc ctagtctggg 180
gagtcctcca taaagttttg catggagcaa acaaacagga ttaaactagg tttggttcct 240
tcagccctct aaaagcatag ggcttagcct gcaggcttcc ttgggctttc tctgtgtgtg 300
tagttttgta aacactatag catctgttaa gatccagtgt ccatggaaac cttcccacat 360
gccgtgactc tggactatat cagtttttgg aaagcagggt tcctctgcct gctaacaagc 420
ccacgtggac cagtctgaat gtctttcctt tacacctatg tttttaata gtcaaacttc 480
aagaaacaat ctaaacaagt ttctgttgca tatgtgtttg tgaacttgta tttgtattta 540
gtaggcttct atattgcatt taacttgttt ttgtaactcc tgattcttcc ttttcggata 600
ctattgatga ataaagaaat t                                     621

```

<210> 210

<211> 533

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 20, 21, 61

<223> n = A,T,C or G

<400> 210

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cgcccttgggg agccggcggn ngagtccggg acgtggagac cgggggtccc ggcagccggg 60
nggcccgcgg gccaggggtg gggatgcacc gccgcggggg gggagctggc gccatcgcca 120
agaagaaact tgcagaggcc aagtataagg agcgagggac ggtcttggct gaggaccagc 180
tagcccagat gtcaaagcag ttggacatgt tcaagaccaa cctggaggaa tttgccagca 240
aacacaagca ggagatccgg aagaatcctg agttccgtgt gcagttccag gacatgtgtg 300
caaccattgg cgtggatccg ctggcctctg gaaaaggatt ttggtctgag atgctgggcg 360
tgggggactt ctattacgaa ctagggtgtcc aaattatcga agtgtgcctg gcgctgaagc 420
atcggaatgg aggtctgata actttggagg aactacatca acaggtgttg aaggggaagg 480
gcaagttcgc ccaggatgtc agtcaagatg acctgatcag agccatcaag aaa 533

```

<210> 211

<211> 451

<212> DNA

<213> Homo sapiens

<400> 211

```

ttagcttgag ccgagaacga ggcgagaaag ctggagaccg aggagaccgc ctagagcgga 60
gtgaacgggg aggggaccgt ggggaccggc ttgatcgtgc gcggacacct gctaccaagc 120
ggagcttcag caaggaagtg gaggagcgga gtagagaacg gccctcccag cctgaggggc 180
tgcgcaaggc agctagcctc acggaggatc gggaccgtgg gcgggatgcc gtgaagcgag 240
aagctgccct acccccagtg agccccctga aggcggctct ctctgaggag gagttagaga 300
agaaatccaa ggctatcatt gaggaatata tccatctcaa tgacatgaaa gaggcagtcc 360
agtgcgtgca ggagctggcc tcacctcct tgctcttcat ctttgtacgg catggtgtcg 420
agtctacgct ggagcgcagt gccattgctc g                                     451

```

<210> 212

<211> 471

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 54  
 <223> n = A,T,C or G

<400> 212  
 gtgattattc ttgatcaggg agaagatcat ttagatttgt tttgcattcc ttanaatgga 60  
 gggcaacatt ccacagctgc cctggctgtg atgagtgtcc ttgcaggggc cggagtagga 120  
 gcactggggg gggggcggaa ttgggggttac tcgatgtaag ggattccttg ttgttggtgt 180  
 gagatccagt gcagttgtga tttctgtgga tcccagcttg gttccaggaa ttttgtgtga 240  
 ttggcttaaa tccagttttc aatcttcgac agctgggctg gaacgtgaac tcagtagctg 300  
 aacctgtctg acccggtcac gttcttggat cctcagaact ctttgcctct gtcgggggtg 360  
 ggggtgggaac tcacgtgggg agcgggtggc gagaaaatgt aaggattctg gaatacatat 420  
 tccatgggac tttccttccc tctcctgctt cctcttttcc tgctccctaa c 471

<210> 213  
 <211> 511  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 27, 63, 337, 442  
 <223> n = A,T,C or G

<400> 213  
 ctaattagaa acttgctgta cttttntttt tcttttaggg gtcaaggacc ctctttatag 60  
 ctncatttg cctacaataa attattgcag cagtttgcaa tactaaaata ttttttatag 120  
 actttatatt tttccttttg ataaagggat gctgcatagt agagttgggt taattaaact 180  
 atctcagccg tttccctgct tttccttctg ctccatatgc ctcatgtcc ttccaggag 240  
 ctcttttaaa cttaaagttc tacatttcat gctcttagtc aaattctgtt acctttttaa 300  
 taactcttcc cactgcata tttccatttg aattggnggt tctaaattct gaaactgtag 360  
 ttgagataca gctatttaaat atttctggga gatgtgcac cctcttcttt gtggttgcc 420  
 aagggtgttt tgcgtaactg anactccttg atatgcttca gagaatttag gcaaacactg 480  
 gccatggccg tgggagtact gggagtaaaa t 511

<210> 214  
 <211> 521  
 <212> DNA  
 <213> Homo sapiens

<400> 214  
 agcattgcc aataatccct aattttccac taaaaatata atgaaatgat gttaagcttt 60  
 ttgaaaagtt taggttaaac ctactgttg tagattaatg tatttgttgc ttccctttat 120  
 ctggaatgtg gcattagctt ttttatttta accctcttta attcttattc aattccatga 180  
 cttaaaggtg gagagctaaa cactgggatt ttgggataac agactgacag ttttgcataa 240  
 ttataatcgg cattgtacat agaaaggata tggctacctt ttgttaaata tgcactttct 300  
 aaatatcaaa aaagggaat gaagtataaa tcaatttttg tataatctgt ttgaaacatg 360  
 agttttatatt gcttaatat agggctttgc cccttttctg taagtctctt gggatcctgt 420  
 gtagaagctg ttctcattaa acaccaaaca gttaagtcca ttctctggta ctagctacaa 480  
 attcggtttc atattctact taacaattta aataaactga a 521

<210> 215  
 <211> 381  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 17, 20, 60, 61, 365  
 <223> n = A,T,C or G

<400> 215  
 gagcggagag cggaccngtn agagccctga gcagccccac cgccgccgcc ggcctagttn 60  
 ncatcacacc ccgggaggag ccgcagctgc cgcagccggc cccagtcacc atcacccgcaa 120  
 ccatgagcag cgaggccgag acccagcagc cgcccgcgcg cccccccgcc gcccccgccc 180  
 tcagcgccgc cgacaccaag cccggcacta cgggcagcgg cgcagggagc ggtggcccg 240  
 gcggcctcac atcggcggcg cctgccggcg gggacaagaa ggtcatcgca acgaaggttt 300  
 tggaacagt aaaatggttc aatgtaagga acggatatgg ttcatcaac aggaatgaca 360  
 ccaangaaga tgtatttgta c 381

<210> 216  
 <211> 425  
 <212> DNA  
 <213> Homo sapiens

<400> 216  
 ttactaacta ggtcattcaa ggaagtcaag ttaacttaaa catgtcacct aaatgcactt 60  
 gatggtggtg aaatgtccac cttcttaaat ttttaagatg aacttagttc taaagaagat 120  
 aacaggccaa tcctgaaggt actccctggt tgctgcagaa tgtcagatat tttggatggt 180  
 gcataagagt cctatttgcc ccagttaatt caacttttgt ctgcctgttt tgtggactgg 240  
 ctggtctctg tagaactctg tccaaaaagt gcatggaata taacttgtaa agcttccac 300  
 aattgacaat atatatgcat gtgttttaaac caaatccaga aagcttaaac aatagagctg 360  
 cataatagta tttattaaag aatcacaact gtaaacaatga gaataactta aggattctag 420  
 ttttag 425

<210> 217  
 <211> 181  
 <212> DNA  
 <213> Homo sapiens

<400> 217  
 gagaaaccaa atgatagggt gtagagcctg atgactccaa acaaagccat cccccgatt 60  
 cttcctcctt cttctggtgc tacagctcca agggcccttc accttcattg ctgaaatgga 120  
 actttggctt tttcagtgga agaatatggt gaagggttca ttttgttcta gaaaaaaaaa 180  
 a 181

<210> 218  
 <211> 405  
 <212> DNA  
 <213> Homo sapiens

<400> 218  
 caggccttcc agttcactga caaacatggg gaagtgtgcc cagctggctg gaaacctggc 60  
 agtgatacca tcaagcctga tgtccaaaag agcaaagaat atttctccaa gcagaagtga 120  
 gcgctgggct gttttagtgc caggctgcgg tgggcagcca tgagaacaaa acctcttctg 180  
 tatttttttt ttccattagt aaaacacaag acttcagatt cagccgaatt gtggtgtctt 240  
 acaaggcagg cctttcctac aggggggtgga gagaccagcc tttcttctt tggtaggaat 300  
 ggcctgagtt ggcgttgtgg gcaggctact gggttgatg atgtattagt agagcaaccc 360  
 attaattctt tgtagtttgt attaaacttg aactgagaaa aaaaa 405

<210> 219  
 <211> 216  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 207, 210  
 <223> n = A,T,C or G

<400> 219  
 actccaagag ttagggcagc agagtggagc gatttagaaa gaacatttta aaacaatcag 60  
 ttaatttacc atgtaaaatt gctgtaaatg ataatgtgta cagattttct gttcaaatat 120  
 tcaattgtaa acttcttggt aagactgtta cgtttctatt gcttttgat gggaattgc 180  
 aaaaataaaa aggaaagaac cctcttnaan aaaaaa 216

<210> 220  
 <211> 380  
 <212> DNA  
 <213> Homo sapiens

<400> 220  
 cttacaaatt gcccccatgt gtaggggaca cagaaccctt tgagaaaact tagatttttg 60  
 tctgtacaaa gtctttgcct ttttccttct tcatTTTTTT ccagtagatt aaatttgtca 120  
 atttcatctt tgagggaac tgatttagatg gggttggtt gtgttctgat ggagaaaaca 180  
 gcacccaag gactcagaag atgattttta cagttcagaa cagatgtgtg caatattggt 240  
 gcatgtaata atgttgagt gcagtcacaa gtcattgatt ttatcttagt tcttcattac 300  
 tgcattgaaa aggaaaacct gtctgagaaa atgcctgaca gtttaattta aaactatggt 360  
 gtaagtcctt gacaaaaaaa 380

<210> 221  
 <211> 398  
 <212> DNA  
 <213> Homo sapiens

<400> 221  
 ggtagtaag ctgtcgactt tgtaaaaaag ttaaaaatga aaaaaaaagg aaaaatgaat 60  
 tgtatatTTA atgaatgaac atgtacaatt tgccactggg aggagggtcc tttttgttgg 120  
 gtgagtcctgc aagtgaattt cactgatgtt gatattcatt gtgtgtagtt ttatttcggt 180  
 cccagccccg tttcctttta ttttggagct aatgccagct gcgtgtctag ttttgagtgc 240  
 agtaaaatag aatcagcaaa tcaactcttat ttttcatcct tttccggtat tttttgggtt 300  
 gtttctgtgg gagcagtgt caccactct tctgtatat tgccTTTTTg ctggaaaatg 360  
 ttgtatgttg aataaaattt tctataaaaa ttaaaaaa 398

<210> 222  
 <211> 301  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 49, 64  
 <223> n = A,T,C or G

<400> 222  
 ttcgataatt gatctcatgg gctttccctg gaggaaaggt tttttttgnt gtttattttt 60  
 taanaacttg aaacttgtaa actgagatgt ctgtagcttt tttgcccac tgtagtgtat 120  
 gtgaagattt caaaacctga gagcactttt tctttgttta gaattatgag aaaggcacta 180  
 gatgacttta ggatttgcac ttttcccttt attgcctcat ttcttgtgac gccttggttg 240  
 ggagggaaat ctgtttattt tttcctacaa ataaaaagct aagattctat atcgcaaaaa 300  
 a 301

<210> 223  
 <211> 200  
 <212> DNA  
 <213> Homo sapiens

<400> 223  
 gtaagtgcctt aggaagaaac tttgcaaaca tttaatgagg atacactggt cattttttaa 60  
 attccttcac actgtaattt aatgtgtttt atattctttt gtagtaaaac aacataaactc 120  
 agatttctac aggagacagt ggttttattt ggattgtcct ctgtaatagg tttcaataaa 180  
 gctggatgaa cttaaaaaaa 200

<210> 224  
 <211> 385  
 <212> DNA  
 <213> Homo sapiens

<400> 224  
 gaaagggtttg atccggactc aaagaaagca aaggagtgtg agccgccatc tgctggagca 60  
 gctgtaactg caagacctgg acaagagatt cgtcagcgaa ctgcagctca aagaaacctt 120  
 tctccaacac cagcaagccc taaccagggc cctcctccac aagttccagt atctcctgga 180  
 ccaccaaaag acagttctgc ccctgggtgga cccccagaaa ggactgttac tccagcccta 240  
 tcatcaaatg tgttaccaag acatcttgga tcccctgcta cttcagtgcc tggaaatgggt 300  
 aaacagagca cttaatgtta tttacagttt atattgtttt ctctgggttac caataaaaacg 360  
 ggccattttc aggtgggtaaa aaaaa 385

<210> 225  
 <211> 560  
 <212> PRT  
 <213> Homo sapiens

<400> 225  
 Met Glu Cys Leu Tyr Tyr Phe Leu Gly Phe Leu Leu Leu Ala Ala Arg  
 1 5 10 15  
 Leu Pro Leu Asp Ala Ala Lys Arg Phe His Asp Val Leu Gly Asn Glu  
 20 25 30  
 Arg Pro Ser Ala Tyr Met Arg Glu His Asn Gln Leu Asn Gly Trp Ser  
 35 40 45  
 Ser Asp Glu Asn Asp Trp Asn Glu Lys Leu Tyr Pro Val Trp Lys Arg  
 50 55 60  
 Gly Asp Met Arg Trp Lys Asn Ser Trp Lys Gly Gly Arg Val Gln Ala  
 65 70 75 80  
 Val Leu Thr Ser Asp Ser Pro Ala Leu Val Gly Ser Asn Ile Thr Phe  
 85 90 95  
 Ala Val Asn Leu Ile Phe Pro Arg Cys Gln Lys Glu Asp Ala Asn Gly  
 100 105 110

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Asn | Ile | Val | Tyr | Glu | Lys | Asn | Cys | Arg | Asn | Glu | Ala | Gly | Leu | Ser | Ala |
|     |     | 115 |     |     |     |     | 120 |     |     |     |     | 125 |     |     |     |
| Asp | Pro | Tyr | Val | Tyr | Asn | Trp | Thr | Ala | Trp | Ser | Glu | Asp | Ser | Asp | Gly |
|     |     | 130 |     |     |     | 135 |     |     |     |     | 140 |     |     |     |     |
| Glu | Asn | Gly | Thr | Gly | Gln | Ser | His | His | Asn | Val | Phe | Pro | Asp | Gly | Lys |
| 145 |     |     |     |     | 150 |     |     |     |     | 155 |     |     |     |     | 160 |
| Pro | Phe | Pro | His | His | Pro | Gly | Trp | Arg | Arg | Trp | Asn | Phe | Ile | Tyr | Val |
|     |     |     | 165 |     |     |     |     | 170 |     |     |     |     |     | 175 |     |
| Phe | His | Thr | Leu | Gly | Gln | Tyr | Phe | Gln | Lys | Leu | Gly | Arg | Cys | Ser | Val |
|     |     |     | 180 |     |     |     |     | 185 |     |     |     |     | 190 |     |     |
| Arg | Val | Ser | Val | Asn | Thr | Ala | Asn | Val | Thr | Leu | Gly | Pro | Gln | Leu | Met |
|     |     | 195 |     |     |     |     | 200 |     |     |     |     | 205 |     |     |     |
| Glu | Val | Thr | Val | Tyr | Arg | Arg | His | Gly | Arg | Ala | Tyr | Val | Pro | Ile | Ala |
|     | 210 |     |     |     |     | 215 |     |     |     |     | 220 |     |     |     |     |
| Gln | Val | Lys | Asp | Val | Tyr | Val | Val | Thr | Asp | Gln | Ile | Pro | Val | Phe | Val |
| 225 |     |     |     |     | 230 |     |     |     |     | 235 |     |     |     |     | 240 |
| Thr | Met | Phe | Gln | Lys | Asn | Asp | Arg | Asn | Ser | Ser | Asp | Glu | Thr | Phe | Leu |
|     |     |     |     | 245 |     |     |     |     | 250 |     |     |     |     | 255 |     |
| Lys | Asp | Leu | Pro | Ile | Met | Phe | Asp | Val | Leu | Ile | His | Asp | Pro | Ser | His |
|     |     | 260 |     |     |     |     | 265 |     |     |     |     |     | 270 |     |     |
| Phe | Leu | Asn | Tyr | Ser | Thr | Ile | Asn | Tyr | Lys | Trp | Ser | Phe | Gly | Asp | Asn |
|     |     | 275 |     |     |     |     | 280 |     |     |     |     | 285 |     |     |     |
| Thr | Gly | Leu | Phe | Val | Ser | Thr | Asn | His | Thr | Val | Asn | His | Thr | Tyr | Val |
|     | 290 |     |     |     |     | 295 |     |     |     |     | 300 |     |     |     |     |
| Leu | Asn | Gly | Thr | Phe | Ser | Leu | Asn | Leu | Thr | Val | Lys | Ala | Ala | Ala | Pro |
| 305 |     |     |     |     | 310 |     |     |     |     | 315 |     |     |     |     | 320 |
| Gly | Pro | Cys | Pro | Pro | Pro | Pro | Pro | Pro | Pro | Arg | Pro | Ser | Lys | Pro | Thr |
|     |     |     | 325 |     |     |     |     |     | 330 |     |     |     |     | 335 |     |
| Pro | Ser | Leu | Gly | Pro | Ala | Gly | Asp | Asn | Pro | Leu | Glu | Leu | Ser | Arg | Ile |
|     |     | 340 |     |     |     |     | 345 |     |     |     |     |     | 350 |     |     |
| Pro | Asp | Glu | Asn | Cys | Gln | Ile | Asn | Arg | Tyr | Gly | His | Phe | Gln | Ala | Thr |
|     |     | 355 |     |     |     |     | 360 |     |     |     |     | 365 |     |     |     |
| Ile | Thr | Ile | Val | Glu | Gly | Ile | Leu | Glu | Val | Asn | Ile | Ile | Gln | Met | Thr |
|     | 370 |     |     |     |     | 375 |     |     |     |     | 380 |     |     |     |     |
| Asp | Val | Leu | Met | Pro | Val | Pro | Trp | Pro | Glu | Ser | Ser | Leu | Ile | Asp | Phe |
| 385 |     |     |     |     | 390 |     |     |     |     | 395 |     |     |     |     | 400 |
| Val | Val | Thr | Cys | Gln | Gly | Ser | Ile | Pro | Thr | Glu | Val | Cys | Thr | Ile | Ile |
|     |     |     | 405 |     |     |     |     |     | 410 |     |     |     |     | 415 |     |
| Ser | Asp | Pro | Thr | Cys | Glu | Ile | Thr | Gln | Asn | Thr | Val | Cys | Ser | Pro | Val |
|     |     |     | 420 |     |     |     |     | 425 |     |     |     |     | 430 |     |     |
| Asp | Val | Asp | Glu | Met | Cys | Leu | Leu | Thr | Val | Arg | Arg | Thr | Phe | Asn | Gly |
|     |     | 435 |     |     |     |     | 440 |     |     |     |     | 445 |     |     |     |
| Ser | Gly | Thr | Tyr | Cys | Val | Asn | Leu | Thr | Leu | Gly | Asp | Asp | Thr | Ser | Leu |
|     | 450 |     |     |     |     | 455 |     |     |     |     | 460 |     |     |     |     |
| Ala | Leu | Thr | Ser | Thr | Leu | Ile | Ser | Val | Pro | Asp | Arg | Asp | Pro | Ala | Ser |
| 465 |     |     |     |     | 470 |     |     |     |     | 475 |     |     |     |     | 480 |
| Pro | Leu | Arg | Met | Ala | Asn | Ser | Ala | Leu | Ile | Ser | Val | Gly | Cys | Leu | Ala |
|     |     |     | 485 |     |     |     |     |     | 490 |     |     |     |     | 495 |     |
| Ile | Phe | Val | Thr | Val | Ile | Ser | Leu | Leu | Val | Tyr | Lys | Lys | His | Lys | Glu |
|     |     | 500 |     |     |     |     |     | 505 |     |     |     |     | 510 |     |     |
| Tyr | Asn | Pro | Ile | Glu | Asn | Ser | Pro | Gly | Asn | Val | Val | Arg | Ser | Lys | Gly |
|     | 515 |     |     |     |     |     | 520 |     |     |     |     | 525 |     |     |     |
| Leu | Ser | Val | Phe | Leu | Asn | Arg | Ala | Lys | Ala | Val | Phe | Phe | Pro | Gly | Asn |
|     | 530 |     |     |     |     | 535 |     |     |     |     | 540 |     |     |     |     |

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 Ala Val Gly Leu Ser Ala Glu Ala Leu Thr Ile Gln Gln Tyr Ala Asp  
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 Val Leu Ser Lys Ala Leu Gly Lys Glu Val Arg Asp Ala Lys Ile Thr  
 85 90 95  
 Pro Glu Ala Phe Glu Lys Leu Gly Phe Pro Ala Ala Lys Glu Ile Ala  
 100 105 110

Asn Met Cys Arg Phe Tyr Glu Met Lys Pro Asp Arg Asp Val Asn Leu  
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| catacactgc  | gacatcgat   | aacggttact  | gtttcacatt  | caccaccctg  | aattgactct  | 4620 |
| cttccggggc  | ctatcatgcc  | ataccgcgaa  | aggttttgcg  | ccattcgatg  | gtgtccggga  | 4680 |
| tctcgacgct  | ctcccttatg  | cgaactcctgc | attaggaagc  | agcccaagtag | taggttgagg  | 4740 |

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| ccggccacgg  | ggcctgccac  | catacccacg  | ccgaaacaag  | cgctcatgag  | cccgaagtgg  | 4860 |
| cgagcccgat  | cttccccatc  | ggtgatgtcg  | gcgatatagg  | cgccagcaac  | cgcacctgtg  | 4920 |
| gcgcgggtga  | tgccggccac  | gatgcgtccg  | gcgtagagga  | tcgagatctc  | gatcccgcg   | 4980 |
| aattaatacg  | actcactata  | ggggaattgt  | gagcggataa  | caattcccct  | ctagaaataa  | 5040 |
| ttttgtttta  | ctttaagaag  | gagatatata  | tatgcagcat  | caccaccatc  | accacggagt  | 5100 |
| acagcttcaa  | gacaatgggt  | ataatggatt  | gctcattgca  | attaatcctc  | aggtacctga  | 5160 |
| gaatcagaac  | ctcatctcaa  | acattaagga  | aatgataact  | gaagcttcat  | tttacctatt  | 5220 |
| taatgctacc  | aagagaagag  | tattttttcag | aaatataaag  | attttaatac  | ctgccacatg  | 5280 |
| gaaagcta    | aataacagca  | aaataaaaaca | agaatcatat  | gaaaaggcaa  | atgtcatagt  | 5340 |
| gactgactgg  | tatggggcac  | atggagatga  | tccatacacc  | ctacaatata  | gaggggtgtg  | 5400 |
| aaaagaggga  | aaatacattc  | atttcacacc  | taatttccta  | ctgaatgata  | acttaacagc  | 5460 |
| tggtacggga  | tcacgaggcc  | gagtgtttgt  | ccatgaatgg  | gcccacctcc  | gttgggggtg  | 5520 |
| gttcgatgag  | tataacaatg  | acaaaccttt  | ctacataaat  | gggcaaaatc  | aaattaaagt  | 5580 |
| gacaagggtg  | tcactctgaca | tcacaggcat  | ttttgtgtgt  | gaaaaaggtc  | cttgccccc   | 5640 |
| agaaaactgt  | attattagta  | agctttttta  | agaaggatgc  | acctttatct  | acaatagcac  | 5700 |
| ccaaaatgca  | actgcatcaa  | taatgttcat  | gcaaagttta  | tcttctgtgg  | ttgaattttg  | 5760 |
| taatgcaagt  | accacaacc   | aagaagcacc  | aaacctacag  | aaccagatgt  | gcagctcag   | 5820 |
| aagtgcattg  | gatgtaatca  | cagactctgc  | tgactttcac  | cacagcttct  | ccatgaacgg  | 5880 |
| gactgagctt  | ccacctctct  | ccacattctc  | gcttgttagag | gctggtgaca  | aagtgggtctg | 5940 |
| tttagtgctg  | gatgtgtcca  | gcaagatggc  | agaggctgac  | agactccttc  | aactacaaca  | 6000 |
| agccgcagaa  | ttttatttga  | tgacagattgt | tgaaattcat  | accttcgtgg  | gcattgccag  | 6060 |
| tttcgacagc  | aaaggagaga  | tcagagccca  | gctacaccaa  | attaacagca  | atgatgatcg  | 6120 |
| aaagttgctg  | gtttcatatc  | tgcccaccac  | tgtatcagct  | aaaacagaca  | tcagcatttg  | 6180 |
| ttcagggctt  | aagaaaggat  | ttgaggtggt  | tgaaaaactg  | aatggaaaag  | cttatggctc  | 6240 |
| tgtgatgata  | ttagtgaacca | gcggagatga  | taagcttctt  | ggcaattgct  | taccactgtg  | 6300 |
| gctcagcagt  | ggttcaacaa  | ttcactccat  | tgccctgggt  | tcactctgcag | ccccaaatct  | 6360 |
| ggaggaatta  | tcacgtctta  | caggaggttt  | aaagtctttt  | gttccagata  | tatcaaactc  | 6420 |
| caatagcatg  | attgatgctt  | tcagtagaat  | ttcctctgga  | actggagaca  | ttttccagca  | 6480 |
| acataattcag | cttgaaaagta | caggtgaaaa  | tgtcaaacct  | caccatcaat  | tgaaaaacac  | 6540 |
| agtgactgtg  | gataatactg  | tgggcaacga  | cactatgttt  | ctagttacgt  | ggcaggccag  | 6600 |
| tggtcctcct  | gagattatat  | tatttgatcc  | tgatggacga  | aaataactaca | caaataattt  | 6660 |
| tatcaccaat  | ctaacttttc  | ggacagctag  | tctttggatt  | ccaggaacag  | ctaagcctgg  | 6720 |
| gcactggact  | tacacctga   | acaataccca  | tcattctctg  | caagccctga  | aagtgacagt  | 6780 |
| gacctctcgc  | gectccaact  | cagctgtgcc  | cccagccact  | gtggaagcct  | ttgtggaaag  | 6840 |
| agacagcctc  | cattttcctc  | atcctgtgat  | gatttatgcc  | aatgtgaaac  | agggatttta  | 6900 |
| tcccattctt  | aatgccactg  | tcactgccac  | agttgagcca  | gagactggag  | atcctgttac  | 6960 |
| gctgagactc  | cttgatgatg  | gagcaggtgc  | tgatgttata  | aaaaatgatg  | gaatttactc  | 7020 |
| gaggtatttt  | ttctcctttg  | ctgcaaatag  | tagatatagc  | ttgaaagtgc  | atgtcaatca  | 7080 |
| ctctcccagc  | ataagcaccc  | cagcccactc  | tattccaggg  | agtcatgcta  | tgtatgtacc  | 7140 |
| aggttacaca  | gcaaacggta  | atattcagat  | gaatgctcca  | aggaaatcag  | taggcagaaa  | 7200 |
| tgaggaggag  | cgaaagtggg  | gcttttagccg | agtcagctca  | ggaggctcct  | tttcagtgtc  | 7260 |
| gggagtcca   | gctggccccc  | accctgatgt  | gtttccacca  | tgcaaaatta  | ttgacctgga  | 7320 |
| agctgtaaaa  | gtagaagagg  | aattgaccct  | atcttgagca  | gcacctggag  | aagactttga  | 7380 |
| tcagggccag  | gctacaagct  | atgaaataag  | aatgagtaaa  | agtctacaga  | atatccaaga  | 7440 |
| tgactttaac  | aatgctattt  | tagtaataac  | atcaaaagca  | aatcctcagc  | aagctggcat  | 7500 |
| caggggagata | tttacgttct  | caccccaaat  | ttccacgaat  | ggacctgaac  | atcagccaaa  | 7560 |
| tgagaaaaca  | catgaaagcc  | acagaattta  | tggttgcaata | cgagcaatgg  | ataggaactc  | 7620 |
| cttacagtct  | gctgtatcta  | acattgcccc  | ggcgccctctg | tttattcccc  | ccaattctga  | 7680 |
| tcctgtacct  | gccagagatt  | atcttatatt  | gaaaggagtt  | ttaacagcaa  | tggttttgat  | 7740 |
| aggaatcatt  | tgcccttatta | tagttgtgac  | acatcatact  | ttaaagcagga | aaaagagagc  | 7800 |
| agacaagaaa  | gagaatggaa  | caaaattatt  | ataatgaatt  | ctgcagatat  | ccatcacact  | 7860 |
| ggcgcccgct  | cgagcaccac  | caccaccacc  | actgagatcc  | ggctgctaac  | aaagcccga   | 7920 |
| aggaagctga  | gttggctgct  | gccaccgctg  | agcaataact  | agcataacct  | cttggggcct  | 7980 |

ctaaacgggt cttgaggggt tttttgctga aaggaggaac tatatccgga t

8031

<210> 255

<211> 401

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 9, 67, 247, 275, 277, 397

<223> n = A,T,C or G

<400> 255

```
gtggccagng actagaaggc gaggcgccgc gggaccatgg cggcggcggc ggacgagcgg 60
agtccanagg acggagaaga cgaggaagag gaggagcagt tggttctggg ggaattatca 120
ggaattattg attcagactt cctctcaaaa tgtgaaaata aatgcaagg tttgggcatt 180
gacactgaga ggcccattct gcaagtggac agctgtgtct ttgctgggga gtatgaagac 240
actctangga cctgtgttat atttgaagaa aatgntnaac atgctgatac agaaggcaat 300
aataaaacag tgctaaaata taaatgccat acaatgaaga agctcagcat gacaagaact 360
ctcctgacag agaagaagga aggagaagaa aacatangtg g 401
```

<210> 256

<211> 401

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 7, 37, 51, 79, 96, 98, 103, 104, 107, 116, 167, 181, 183,  
194, 206, 276, 303, 307, 308, 310, 323, 332, 341, 353, 374,  
376

<223> n = A,T,C or G

<400> 256

```
tggtagncct gggatgggga accgcggtgg ctcccgngga ggtttcggca ntggcatccg 60
gggccggggg cgcggccgng gacggggccg gggccnangc cgngganctc gcggangcaa 120
ggccgaggat aaggagtggg tgcccgtcac caacttgggc cgcttgacca aggacatgaa 180
nancagccc ctgnaggaga tctatntctt ctccctgcc ccattaagga atcaagagat 240
catttgattt ctccctgggg gcctctctca aggatnaggt ttttgaagat tatgccagt 300
canaaannan accccgttgc ccngtccatc tncacccaac ncttccaagg gcnatttttg 360
tttaggcctc attncngggg ggaaccttaa cccaatttgg g 401
```

<210> 257

<211> 401

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 382, 387

<223> n = A,T,C or G



&lt;400&gt; 257

```

atgtatgtaa aacacttcat aaaatgtaaa gggctataac aaatatgtta taaagtgatt 60
ctctcagccc tgaggtatac agaatcattt gcctcagact gctgttggat tttaaaattt 120
ttaaaatatac tgctaagtaa tttgctatgt cttctccac actatcaata tgcctgcttc 180
taacaggctc cccactttct tttaatgtgc tgttatgagc tttggacatg agataaccgt 240
gcctgttcag agtgtctaca gtaagagctg gacaaactct ggaggacac agtctttgag 300
acagctcttt tggttgcttt ccacttttct gaaaggttca cagtaacctt ctagataata 360
gaaactccca gttaaagcct angctancaa ttttttttag t 401

```

&lt;210&gt; 258

&lt;211&gt; 401

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 258

```

ggagcgctag gtcgggtgtac gaccgagatt aggggtgctg ccagctccgg gaggccgcgg 60
tgagggggccg ggcccaagct gccgaccga gccgatcgtc agggtcgcca gcgcctcagc 120
tctgtggagg agcagcagta gtcggagggt gcaggatatt agaaatggct actccccagt 180
caattttcat ctttgcaatc tgcattttta tgataacaga attaatctg gcctcaaaaa 240
gctactatga tatcttaggt gtgccaaaat cggcatcaga gcgccaaatc aagaaggcct 300
ttcacaagtt ggcatgaag taccaccctg aaaaaataa gaccagatg ctgaagcaaa 360
attcagagag attgcagaag catatgaaac actctcagat g 401

```

&lt;210&gt; 259

&lt;211&gt; 401

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 259

```

attgggtttg gagggaggat gatgacagag gaatgccctt tggccatcac ggttttgatt 60
ctccagaata ttgtgggttt gatcatcaat gcagtcattg taggctgcat tttcatgaaa 120
acagctcagg ctacagaag ggcagaaact ttgattttca gccgccatgc tgtgattgcc 180
gtccgaaaatg gcaagctgtg cttcatgttc cgagtgggtg acctgaggaa aagcatgac 240
attagtgcct ctgtgcgcat ccagggtgtc aagaaaacaa ctacacctga aggggagggtg 300
gttcctattc accaactgga cattcctgtt gataacccaa tcgagagcaa taacattttt 360
ctggtggccc ctttgatcat ctgccacgtg attgacaagc g 401

```

&lt;210&gt; 260

&lt;211&gt; 363

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

<222> 7, 9, 19, 41, 63, 73, 106, 111, 113, 116, 119, 156, 158,  
162, 187, 247, 288, 289, 290, 292, 298, 299, 300, 340

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 260

```

aggaganang gagggggana tgaataggga tggagaggga natagtggat gagcaggcca 60
caggagagg aancagaaag gagaggcaag acaggagagac acacancaca nangangana 120
caggtggggg ctgggggtggg gcatggagag cttttnangt cccccaggcc accctgctct 180
cgctggngctg ttgaaaccca ctccatggct tcttgccact gcagttgggc ccagggtgg 240
cttatnctg gaatgcaagt ggctgtggct tggagcctcc cctctggnnn anggaaannn 300

```

```
attgctccct tatctgcttg gaatatctga gtttttccan cccggaaata aaacacacac 360
aca 363
```

```
<210> 261
<211> 401
<212> DNA
<213> Homo sapiens
```

```
<220>
<221> misc_feature
<222> 114, 152
<223> n = A,T,C or G
```

```
<400> 261
cggctctccg ccgctctccc ggggtttcgg ggcacttggg tcccacagtc tggctcctgct 60
tcaccttccc ctgacctgag tagtcgccat ggcacagggt ctcagaggca ctgngactga 120
cttccctgga tttgatgagc gggctgatgc anaaactctt cggaaaggcta tgaaaggctt 180
gggcacagat gaggagagca tcctgactct gttgacatcc cgaagtaatg ctcagcgcca 240
ggaaatctct gcagctttta agactctgtt tggcagggat cttctggatg acctgaaatc 300
agaactaact ggaaaatttg aaaaattaat tgtggctctg atgaaaccct ctcggtttta 360
tgatgcttat gaactgaaac atgccttgaa gggagctgga a 401
```

```
<210> 262
<211> 401
<212> DNA
<213> Homo sapiens
```

```
<220>
<221> misc_feature
<222> 7, 26, 258, 305, 358, 373, 374, 378
<223> n = A,T,C or G
```

```
<400> 262
agtctanaac atttctaata ttttgnctt tcatatatca aaggagatta tgtgaaacta 60
tttttaata ctgtaaagtg acatatagtt ataagatata tttctgtaca gtagagaaag 120
agtttataac atgaagaata ttgtaccatt atacattttc attctcgatc tcataagaaa 180
ttcaaaagaa taatgataga ggtgaaaata tgtttacttt ctctaaatca agcctagtgt 240
tcaactcaaa aattatgntg catagtttta ttttgaattt aggttttggg actacttttt 300
tccancttca atgagaaaaa aaaatctaca actcaggagt tactacagaa gttctaanta 360
tttttttgct aannagcnaa aaatataaac atatgaaaat g 401
```

```
<210> 263
<211> 401
<212> DNA
<213> Homo sapiens
```

```
<220>
<221> misc_feature
<222> 232, 290, 304, 326, 383
<223> n = A,T,C or G
```

```
<400> 263
ctgtccgacc aagagaggcc ggccgagccc gaggcttggg cttttgcttt ctggcggagg 60
gatctgcggc ggtttaggag gcggcgctga tcctgggagg aagaggcagc tacggcggcg 120
```

```

gcgggcggtgg cggctagggc ggcgggcgaat aaagggggccg ccgcccgggtg atgcgggtgac 180
cactgcggca ggcccaggag ctgagtgggc cccggccctc agcccgtccc gncggaccgc 240
ctttccctcaa ctctccatct tctcctgccg accgagatcg ccgaggcggn ctcaggctcc 300
ctanccctt ccccgctcct tcccncccc cgccccgcc ccggggggccg ccgccaccgc 360
cctcccacca tggtcttgaa ganaatccac aaggaattga a 401

```

<210> 264

<211> 401

<212> DNA

<213> Homo sapiens

<400> 264

```

aacaccagcc actccaggac ccctgaaggc ctctaccagg tcaccagtgt tctgcgccta 60
aagccacccc ctggcagaaa cttcagctgt gtgttctgga atactcacgt gagggaaactt 120
actttggcca gcattgacct tcaaagtcag atggaaacca ggaccatcc aacttggtg 180
cttcacattt tcatccctc ctgcatcatt gctttcattt tcatagccac agtgatagcc 240
ctaagaaaaa aactctgtca aaagctgtat tcttcaaaa acacaacaaa aagacctgtc 300
accacaacaa agagggaagt gaacagtgtc gtgaatctga acctgtggtc ttgggagcca 360
gggtgacctg atatgacatc taaagaagct tctggactct g 401

```

<210> 265

<211> 271

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 59

<223> n = A,T,C or G

<400> 265

```

gccatttct gtggacatgg gcagagcgct gctgccagtt cctggtagcc ttgaccacna 60
cgctgggggg tctttgtgat ggtcatgggt ctcatctgca cttgggggtg tgggattcaa 120
gttagaagtt tctagatctg gccgggcgca gtggctcaca cctgtaatcc cagcacttta 180
ggaggctgag gcaggcggat catgaggtca ggagatcgag accgtcctgg ctaacacagt 240
gaaaccccg ctctactaaa aatacaaaaa a 271

```

<210> 266

<211> 401

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 45

<223> n = A,T,C or G

<400> 266

```

attcataaat ttagctgaaa gatactgatt caatttgtat acagngaata taaatgagac 60
gacagcaaaa ttttcatgaa atgtaaaata tttttatagt ttgttcatac tatatgaggt 120
tctattttta atgactttct ggatttttaa aaatttcttt aaatacaatc atttttgtaa 180
tatttatttt atgcttatga tctagataat tgcagaatat cattttatct gactctgtct 240
tcataagaga gctgtggccg aattttgaac atctgttata gggagtgatc aaattagaag 300
gcaatgtgga aaaacaattc tgggaaagat ttctttatat gaagtcctg ccactagcca 360

```

gccatcctaa ttgatgaaag ttatctgttc acaggcctgc a 401

<210> 267

<211> 401

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 116, 247, 277, 296, 307, 313, 322, 323, 336, 342, 355, 365, 377, 378, 397

<223> n = A,T,C or G

<400> 267

gaagaggcat cacctgatcc cggagacctt tggagttaag aggcggcgga agcgagggcc 60  
tgtggagtcg gatcctcttc ggggtgagcc aggttcggcg cgcgcggctg tctcanaact 120  
catgcagctg ttcccgcgag gcctgtttga ggacgcgctg ccgcccacg tgctgaggag 180  
ccaggtgtac agccttgtgc ctgacaggac cgtggccgac cggcagctga aggagcttca 240  
agagcanggg gagacaaaat cgtccagctg ggcttcnact tggatgcca tggaanttat 300  
tctttcnctt ganggactta cnnnggaccc aagaanccct tncaaggggc ccttngtgga 360  
tgggncccga aaccccnnta tttgcccttg ggggggncca a 401

<210> 268

<211> 223

<212> DNA

<213> Homo sapiens

<400> 268

tgcctatgtt ggccaggctg gtcttgaact cctgacttta agtgatccac ccgcctcaac 60  
ctcccaaagt gctgggatta cagggtgtgag ccaccgcgcc tggcctgata catactttta 120  
gaatcaagta gtcacgcact ttttctgttc atttttctaa aaagtaaata taaaaatgtt 180  
ttgttttttg ttttttttgt ttgtttgttt ctgttttttt ttt 223

<210> 269

<211> 401

<212> DNA

<213> Homo sapiens

<400> 269

actatgtaaa ccacattgta ctttttttta ctttggcaac aaatatttat acatacaaga 60  
tgctagttca tttgaatatt tctcccaact tatccaagga tctccagctc taacaaaatg 120  
gtttattttt atttaaagt caatagttgt tttttaaaat ccaaatacaga ggtgcaggcc 180  
accagttaaa tgccgtctat caggttttgt gccttaagag actacagagt caaagctcat 240  
ttttaaaagga gtaggacaaa gttgtcacag gtttttgttg ttgtttttat tgccccaaa 300  
attacatggt aatttccatt tatatcaggg attctattta cttgaagact gtgaagttgc 360  
cattttgtct cattgttttc tttgacataa ctaggatcca t 401

<210> 270

<211> 401

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 240, 382

<223> n = A,T,C or G

<400> 270

```

tggctgttga ttcacctcag cactgcttgg tatctgcacc ctacctctct ttagaggctg 60
ccttgtcaac tgaaaaatgc acctgacttc gagcaagact ctttccttag gttctggatc 120
tgtttgagcc ccatggcact gagctggaat ctgagggtct tgttccaagg atgtgatgat 180
gtgggagaat gttctttgaa agagcagaaa tccagtctgc atggaaacag cctgtagagn 240
agaagtttcc agtgataagt gttcactgtt ctaaggaggt acaccacagc tacctgaatt 300
ttcccaaaat gagtgcttct gtgcgttaca actggccttt gtacttgact gtgatgactt 360
tgttttttct tttcaattct anatgaacat gggaaaaaat g 401

```

<210> 271

<211> 329

<212> DNA

<213> Homo sapiens

<400> 271

```

ccacagcctc caagtcaggt ggggtggagt cccagagctg cacagggttt ggcccaagtt 60
tctaaggagag gcacttcctc ccctcgccca tcagtgccag cccctgctgg ctggtgcctg 120
agccctcag acagccccct gcccgcagg cctgccttct cagggaactc tgcggggcct 180
gaggcaagcc atggagttag acccaggagc cggacacttc tcaggaaatg gcttttccca 240
acccccagcc cccaccgggt ggttcttctt gttctgtgac tgtgtatagt gccaccacag 300
cttatggcat ctcataggag acaaaaaaa 329

```

<210> 272

<211> 401

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 1, 7, 12, 21, 61, 62, 66, 72, 78, 88, 90, 92, 98, 117, 119, 128, 130, 134, 142, 144, 151, 159, 162, 164, 168, 169, 177, 184, 185, 188, 194, 202, 204, 209, 213, 218, 223, 231, 260, 272, 299, 300, 306, 321, 322, 323, 331, 335, 336, 338

<223> n = A,T,C or G

<221> misc\_feature

<222> 341, 342, 343, 345, 346, 351, 358, 360, 362, 363, 387, 390, 392

<223> n = A,T,C or G

<400> 272

```

nggctgntaa cntcggaggt nacttcctgg actatcctgg agacccccctc cgcttccacg 60
nncatnatat cntcatngc tggggccntn angacacnat cccactccaa cacctgngng 120
atgctggncn cctnggaacc ancntcagaa ngaccctgnt cntntgtntt ccgcaanctg 180
aagnnaangc gggntacacc tnentgcant ggnccacnct gcngggaact ntacacacct 240
acgggatgtg gctgcgccan gagccaagag cntttctgga tgattcccca gcctcttggn 300
agggantcta caacattgct nnntaccttt ntcennngc nnntnntgga ntacaggngn 360
tnntaact acatcttttt tactgncncc tncttgggtg g 401

```

<210> 273

<211> 401

<212> DNA  
 <213> Homo sapiens  
 <220>  
 <221> misc\_feature  
 <222> 399  
 <223> n = A,T,C or G

<400> 273  
 cagcaccatg aagatcaaga tcatcgcacc cccagagcgc aagtactcgg tgtggatcgg 60  
 tggctccatc ctggcctcac tgtccacctt ccagcagatg tggattagca agcaggagta 120  
 cgacgagtcg ggcccctcca tcgtccaccg caaatgcttc taaacggact cagcagatgc 180  
 gtagcatttg ctgcatgggt taattgagaa tagaaatttg cccctggcaa atgcacacac 240  
 ctcatgctag cctcacgaaa ctggaataag ccttcgaaaa gaaattgtcc ttgaagcttg 300  
 tatctgatat cagcaactgga ttgtagaact tgttgctgat tttgaccttg tattgaagtt 360  
 aactgttccc ctgtgtatta acgtgtcagg gctgagtgnt c 401

<210> 274  
 <211> 401  
 <212> DNA  
 <213> Homo sapiens

<400> 274  
 ccacccacac ccaccgcgcc ctctgttcgcc tcttctccgg gagccagtcc gcgccaccgc 60  
 cgccgccag gccatcgcca cctccgcag ccatgtccac cagggtccgtg tctctgtcct 120  
 cctaccgcag gatgttcggc ggcccgggca ccgcgagccg gccgagctcc agccggagct 180  
 acgtgactac gtccacccgc acctacagcc tgggcagcgc gctgcgcccc agcaccagcc 240  
 gcagcctcta cgctctgtcc ccgggcggcg tgtatgccac gcgtctctct gccgtgcgcc 300  
 tgcggagcag cgtgcccggg gtgcggctcc tgcaggactc ggtggacttc tcgtctggccg 360  
 acgccatcaa caccgagttc aagaacaccc gcaccaacga g 401

<210> 275  
 <211> 401  
 <212> DNA  
 <213> Homo sapiens

<400> 275  
 ccacttccac cactttgtgg agcagtgcct tcagcgcaac ccggatgccca ggtatccctg 60  
 ctggcctggg cctgggcttc gggagagcag aggggtgctca ggagggtaag gccaggggtg 120  
 gaagggactt acctcccaaa ggttctgcag gggaatctgg agctacacac aggagggatc 180  
 agctcctggg tgtgtcagag gccagcctgg ggagctctgg ccactgtctc ccatgagctg 240  
 agggagaggg agaggggacc cgaggctgag gcataagtgg caggatttcg ggaagctggg 300  
 gacacggcag tgatgctgcg gtctctctct ccctttccct ccaggcccag tgccagcacc 360  
 ctctgaacc actctttctt caagcagatc aagcgacgtg c 401

<210> 276  
 <211> 401  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 11  
 <223> n = A,T,C or G

```

<400> 276
tctgatattg ntacccttga gccacctaag ttagaagaaa ttggaaatca agaagttgtc 60
attgttgaag aagcacagag ttcagaagac tttaacatgg gctcttcctc tagcagccag 120
tatactttct gtcagccaga aactgtatct tcattctcagc ctagtgtatga tgaatcaagt 180
agtgtatgaaa ccagtaatca gccagtcctt gccttttagac gacgccgtgc taggaagaag 240
accgtttctg cttcagaatc tgaagaccgg ctagtgtgtg aacaagaaac tgaaccttct 300
aaggagttga gtaaaccgtc gttcagtagt ggtctcaata agtgtgttat acttgctttg 360
gtgattgcaa tcagcatggg atttggccat ttctatggca c 401

```

<210> 277

<211> 401

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 227, 333

<223> n = A,T,C or G

<400> 277

```

aactttggca acatatctca gcaaaaacta cagctatggt attcatgcc aataaaaagc 60
tgtgcagagg agtggctgca atgaggtcac aacggtggtg gatgtaaaag agatcttcaa 120
gtcctcatca cccatccctc gaactcaagt cccgctcatt acaaattctt cttgccagt 180
tccacacatc ctgccccatc aagatgttct catcatgtgt tacgagnggc gctcaaggat 240
gatgcttctt gaaaattgct tagttgaaaa atggagagat cagcttagta aaagatccat 300
acagtgggaa gagaggctgc aggaacagcg ganaacagtt caggacaaga agaaaacagc 360
cgggcgcacc agtcgtagta atccccccaa accaaaaggga a 401

```

<210> 278

<211> 401

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 322, 354

<223> n = A,T,C or G

<400> 278

```

aatgagtgtg agaccacaaa tgaatgccgg gaggatgaaa tgtgttggaa ttatcatggc 60
ggcttccggt gttatccacg aaatccttgt caagatccct acattctaac accagagaac 120
cgatgtgttt gccagtcctc aaatgccatg tgccgagaaac tgccccagtc aatagtctac 180
aaatacatga gcatccgacg tgataggtct gtgccatcag acatcttcca gatacaggcc 240
acaactatct atgccaacac catcaatact ttctcgatta aatctggaaa tgaaaatgga 300
gagtctacct acgacaacaa anccctgtaa gtgcaatgct tgtgctcgtg aagncattat 360
caggaccaag agaacatata gtggacctgg agatgctgac a 401

```

<210> 279

<211> 401

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature  
 <222> 30, 35, 81, 88, 180, 212, 378, 384, 391  
 <223> n = A,T,C or G

<400> 279  
 aaattattgc ctctgataca tacctaagtn aacanaacat taatacctaa gtaaacataa 60  
 cattacttgg agggttgcag nttctaantg aaactgtatt tgaaactttt aagtatactt 120  
 taggaaacaa gcatgaacgg cagtctagaa taccagaaac atctacttgg gtagcttggn 180  
 gccattatcc tgtggaatct gatatgtctg gnagcatgtc attgatggga catgaagaca 240  
 tctttgaaa tgatgagatt atttcctgtg ttaaaaaaaa aaaaaatctt aaattcctac 300  
 aatgtgaaac tgaaactaat aattttgatc ctgatgtatg ggacagcgta tctgtaccag 360  
 gctctaaata acaaaagnta gggngacaag nacatgttcc t 401

<210> 280  
 <211> 326  
 <212> DNA  
 <213> Homo sapiens

<400> 280  
 gaagtggaat tgtataattc aattcgataa ttgatctcat gggctttccc tggaggaaaag 60  
 gttttttttg ttgttttttt tttaagaact tgaaacttgt aaactgagat gtctgtagct 120  
 tttttgcca tctgtagtgt atgtgaagat ttcaaaacct gagagcactt tttctttgtt 180  
 tagaattatg agaaaggcac tagatgactt taggatttgc atttttccct ttattgcctc 240  
 atttcttgtg acgccttggt ggggagggaa atctgtttat tttttcctac aaataaaaaag 300  
 ctaagattct atatcgcaaa aaaaaa 326

<210> 281  
 <211> 374  
 <212> DNA  
 <213> Homo sapiens

<400> 281  
 caacgcgttt gcaaatatct ccctggtagc ctacttcctt acccccgaat attggttaaga 60  
 tcgagcaatg gcttcaggac atgggttctc ttctcctgtg atcattcaag tgctcactgc 120  
 atgaagactg gcttgtctca gtgtttcaac ctcaccaggg ctgtctcttg gtccacacct 180  
 cgctccctgt tagtgccgta tgacagcccc catcaaatga ccttggccaa gtcacggttt 240  
 ctctgtggtc aaggttgggt ggctgattgg tggaaagtag ggtggaccaa aggaggccac 300  
 gtgagcagtc agcaccagtt ctgcaccagc agcgccctccg tctagtggg tgttcctgtt 360  
 tctcctggcc ctgg 374

<210> 282  
 <211> 404  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 26, 27, 51, 137, 180, 222  
 <223> n = A,T,C or G

<400> 282  
 agtgtgtgg aattcccgca tcctanncg cgactcacac aaggcagagt ngccatggag 60  
 aaaattccag tgtcagcatt ctgtctcctt gtggccctct cctacactct ggccagagat 120  
 accacagtca aacctgnagc caaaaaggac acaaaggact ctcgacccaa actgccccan 180



```

accctctcca gaggttgagg tgaccaactc atctggactc anacatatga agaagctcta 240
tataaatcca agacaagcaa caaacccttg atgattattc atcacttga tgagtgccca 300
cacagtcaag ctttaaagaa agtgtttgct gaaaataaag aaatccagaa attggcagag 360
cagtttgctc tcctcaatct ggtttatgaa acaactgaca aaca 404

```

```

<210> 283
<211> 184
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 26
<223> n = A,T,C or G

```

```

<400> 283
agtgtggtgg aattcacttg cttaanttgt gggcaaaaga gaaaaagaag gattgatcag 60
agcattgtgc aatacagttt cattaactcc ttccctcgct cccccaaaaa tttgaatttt 120
tttttcaaca ctcttacacc tgttatggaa aatgtcaacc tttgtaagaa aacccaaaata 180
aaaa 184

```

```

<210> 284
<211> 421
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 147, 149
<223> n = A,T,C or G

```

```

<400> 284
ctattaatcc tgccacaata tttttaatta cgtacaaaga tctgacatgt caccagggga 60
cccatttcac ccactgctct gtttgccgc cagtcttttg tctctctctt cagcaatggg 120
gaggcgata ccctttcctc ggggaanana aatccatggg ttggtgccct tgccaataac 180
aaaaatggtg gaaagtcgag tggcaaagct gttgccattg gcattcttca cgtgaaccac 240
gtcaaaagat ccagggtgcc tctctctggt ggtgatcaca ccaattcttc ctagggttagc 300
acctccagtc accatacaca gggtaccagt gtcgaacttg atgaaatcag taatcttgcc 360
agtctctaaa tcaatctgaa tgggtatcatt caccttgatg aggggatcgg ggtagcggat 420
g 421

```

```

<210> 285
<211> 361
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 34, 188
<223> n = A,T,C or G

```

```

<400> 285
ctgggtggta actctttatt tcattgtccg gaanaaagat gggagtggga acagggtgga 60
cactgtgcag gcttcagctt ccactccggg caggattcag gctatctggg accgcaggga 120

```

```

ctgccagggtg cacagccctg gctcccgagg caggcaggca aggtgacggg actggaagcc 180
cttttcanag ccttggagga gctgggtccgt ccacaagcaa tgagtgccac tctgcagttt 240
gcaggggatg gataaacagg gaaacactgt gcattcctca cagccaacag tgtaggtctt 300
ggtgaagccc cggcgctgag ctaagctcag gctgttccag ggagccacga aactgcagggt 360
a 361

```

```

<210> 286
<211> 336
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 40, 68, 75, 127, 262
<223> n = A,T,C or G

```

```

<400> 286
tttgagtggc agcgccctta tttgtggggg ccttcaaggc agggtcgtgg ggggcagcgg 60
ggaggaanag ccganaaaact gtgtgaccgg ggcctcagggt ggtgggcatt gggggctcct 120
cttgcanaatg cccattggca tcaccgggtgc agccattggt ggcagcgggt accggtcctt 180
tcttgttcaa catagggtag gtggcagcca cgggtccaac tcgcttgagg ctgggccctg 240
ggcgctccat tttgtgttcc angagcatgt ggttctgtgg cgggagcccc acgcaggccc 300
tgaggatggt ctcgatgcag ctgcgctggc ggaaaa 336

```

```

<210> 287
<211> 301
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 15, 33, 44, 53, 76, 83, 107, 117, 154, 166, 192, 194, 207,
215, 241, 246
<223> n = A,T,C or G

```

```

<400> 287
tgggtaccaaa atttntttat ttgaaggaat ggnacaaatc aaanaactta agnggatggt 60
ttggtacaac ttatanaaaa ggnaaaggaa accccaacat gcatgcnctg ctttgngnac 120
cagggaagtc accccacggc tatggggaaa ttancccgag gcttancttt cattatcact 180
gtctcccagg gngngcttgt caaaaanata ttccnccaag ccaaattcgg gcgctcccat 240
nttgcnaaag ttggtcacgt ggtcacccaa ttctttgatg gctttcacct gctcattcag 300
g 301

```

```

<210> 288
<211> 358
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 39, 143, 226
<223> n = A,T,C or G

```

```

<400> 288

```

```

aagtttttaa actttttatt tgcataattaa aaaaattgng cattccaata attaaaatca 60
tttgaacaaa aaaaaaaatg gcaactctgat taaactgcat tacagcctgc aggacacctt 120
gggccagctt gggttttactc tanattttcac tgtcgtccca ccccaacttct tccaccccac 180
ttcttctctc accaacaatgc aagttctttc cttccctgcc agccanatag atagacagat 240
gggaaaggca ggcgcggcct tcgtttgtcag tagttctttg atgtgaaagg ggcagcacag 300
tcattttaaac ttgatccaac ctctttgcat cttacaaagt taaacagcta aaagaagt 358

```

```

<210> 289
<211> 462
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> 87, 141, 182, 220, 269, 327
<223> n = A,T,C or G

```

```

<400> 289
ggcatcagaa atgctgttta tttctctgct gctcccaagc tggctggcct ttgcagagga 60
gcagacaaca gatgcatagt tgggganaaa gggaggacag gttccaggat agaggggtgca 120
ggctgagggg ggaagggtaa naggaaggaa ggccatcctg gatccccaca ttccagtctc 180
anatgaggac aaaggggactc ccaagccccc aaatcatcan aaaaacaccaa ggagcaggag 240
gagcttgagc agggcccagg gagcctcana gccataccag ccactgtcta cttcccatcc 300
tcctctccca ttccctgtct gcttcanacc acctcccagc taagccccag ctccattccc 360
ccaatcctgg cccttgccag cttgacagtc acagtgcctg gaattccacc actgaggctt 420
ctcccagttg gattaggacg tcgcctgttt agcatgctgc cc 462

```

```

<210> 290
<211> 481
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> 44, 57, 122, 158, 304, 325, 352, 405
<223> n = A,T,C or G

```

```

<400> 290
tactttccta aactttatta aagaaaaaag caataagcaa tggnggtaaa tctctanaac 60
atacccaatt ttctgggctt cctcccccca gaatgtgaca ttttgatttc caaacatgcc 120
anaagtgtat ggttcccaac tgtactaaag taggtganaa gctgaagtcc tcaagtgttc 180
atcttccaac ttttcccagt ctgtgggtctg tctttggatc agcaataatt gcctgaacag 240
ctactatggc ttcgttgatt ttgtctgtga gctctctgag ctctctatg tgcagcaatc 300
gcanaatttg agcagcttca ttaanaactg catctcctgt gtcaaaaacca anaatatgtt 360
tgtctaaagc aacaggtaag cctctttttg ttgattttgc cttancaact gcatcctgtg 420
tcaggcgctc ctgaacccaa atccgaattg ccttaagcat taccaggtaa tcatcatgac 480
g 481

```

```

<210> 291
<211> 381
<212> DNA
<213> Homo sapiens

```

```

<220>

```

<221> misc\_feature  
 <222> 79, 166, 187, 208, 219, 315  
 <223> n = A,T,C or G

<400> 291  
 tcatagtaat gtaaaacat ttgtttaatt ctaaatacaa tcactttcac aacagtga 60  
 attagtgact ggttaaggng tgccactgta catatcatca ttttctgact ggggtcagga 120  
 cctggtccta gtccacaagg gtggcaggag gaggggtggag gctaanaaca cagaaaacac 180  
 aaaaaanaaa ggaaagctgc cttggcanaa ggatgaggng gtgagcttgc cgaaggatgg 240  
 tgggaagggg gctccctgtt ggggccgagc caggagtccc aagtcagctc tcctgcctta 300  
 cttagctcct ggcanagggt gagtggggac ctacgaggtt caaaatacaa tggcatttgg 360  
 ccagcctggc ttactaaca g 381

<210> 292  
 <211> 371  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 32, 55, 72, 151, 189, 292  
 <223> n = A,T,C or G

<400> 292  
 gaaaaaataa tccgtttaat tgaaaaacct gnaggatact attccactcc cccanattgag 60  
 gaggctgagg anaccaaacc cctacatcac ctgtagcca cttctgatac tcttcacgag 120  
 gcagcaggca aagacaattc ccaaaacctc naaaaagca attccaaggg ctgctgcagc 180  
 taccaccanc acatttttcc tcagccagcc cccaattctc tccacacagc cctccttatg 240  
 gatcgcttcc tcgttgaaat taatcccaca gccacagta acattaatgc ancaggagtc 300  
 ggggactcgg ttcttcgaca tggaagggat tttctcccaa tctgtgtagt tagcagcccc 360  
 acagcactta a 371

<210> 293  
 <211> 361  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 75, 196, 222  
 <223> n = A,T,C or G

<400> 293  
 gatttaaaag aaaacacttt attgttcagc aattaaaagt tagccaaata tgtatttttc 60  
 tccataattt attngatgt tatcaacatc aagtaaaatg ctcatattca tcatttgctt 120  
 ctgttcattg tttcttgaac acgtcttcaa ttttccttcc aaaatgctgc atgccacac 180  
 tgaggtaacg aagcanaagt atttttaaac atgacagcta anaacattca tctacagcaa 240  
 cctatatgct caatacatgc cgcgtgatcc tagtagtttt ttcacaacct tctacaagtt 300  
 tttggaaaac atctgttatg atgactttca tacaccttca cctcaaaggc tttcttgac 360  
 c 361

<210> 294  
 <211> 391  
 <212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 26, 77, 96, 150, 203, 252, 254, 264, 276

<223> n = A,T,C or G

<400> 294

```
tattttaaaag ttttaattatg attcanaaaaa aatcgagcga ataactttct ctgaaaaaat 60
atattgactc tgtatanacc acagttattg gggganaagg gctggtaggt taaattatcc 120
tattttttat tctgaaaatg atattaatan aaagtcccg ttcagtcctg attataaaga 180
tacatatgcc caaaatggct ganaataaat acaacaggaa atgcaaaaagc tgtaaagcta 240
agggcatgca ananaaaatc tcanaatacc caaagnggca acaaggaacg tttggctgga 300
atttgaagtt atttcagtc tctttgtctt tggtccatg tttcaggatg cgtgtgaact 360
cgatgtaatt gaaattcccc tttttatcaa t 391
```

<210> 295

<211> 343

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 145, 174, 205, 232

<223> n = A,T,C or G

<400> 295

```
ttcttttgtt ttattgataa cagaaactgt gcataattac agatttgatg aggaatctgc 60
aaataataaaa gaatgtgtct actgccagca aaatacaatt attccatgcc ctctcaacat 120
acaaatatag agttcttcac accanatggc tctggtgtaa caaagccatt ttanatgttt 180
aattgtgctt ctacaaaacc ttcanagcat gaggtagttt cttttaccta cnatattttc 240
cacatttcca ttattacact tttagtgagc taaaatcctt ttaacatagc ctgcggatga 300
tctttcacaa aagccaagcc tcatttacaa agggtttatt tct 343
```

<210> 296

<211> 241

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 96, 98, 106, 185

<223> n = A,T,C or G

<400> 296

```
ttcttgataa ttggttggtt ttgtgaaaaa gtttttggtt ttcttctcag tcaactgaat 60
tatttctcta ctttgccctc ctgatgccca catgananaa cttaanataa tttctaacag 120
cttcactttt ggaaaaaaa aaaacctgtt ttccctcatg aaccccagga gttgaaagt 180
gatanatgc tctcaaaatc taaggctctg ttcagcttta cattatgtta cctgacgttt 240
t 241
```

<210> 297

<211> 391

<212> DNA

137

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 12, 130

<223> n = A,T,C or G

<400> 297

```
gttgtggctg anaatgctgg agatgctcag ttctctccct cacaaggtag gccacaaatt 60
cttgggtggg ccttcacatc tggggctctc aggcaccagc catgcctgcc gaggagtgtc 120
gtcaggacan accatgtccg tgctaggccc aggcacagcc caaccactcc tcatccaagt 180
ctctcccagg tttctgggcc cgatgggcaa ggatgacccc tccagtggct ggtacccac 240
catcccacta cccctcacat gctctcactc tccatcaggt cccaatcct ggcttcctc 300
ttcacgaact ctcaaagaaa aggaaggata aaacctaaat aaaccagaca gaagcagctc 360
tggaaaagta caaaaagaca gccagagggtg t 391
```

<210> 298

<211> 321

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 14, 30, 76, 116, 201, 288, 301

<223> n = A,T,C or G

<400> 298

```
caagccaaac tgtntccagc tttattaaan atactttcca taaacaatca tggattttca 60
ggcaggacat gggcanacaa tcgttaacag tataacaaca ctttcaaact ccttnttca 120
atggactacc aaaaatcaaa aagccactat aaaacccaat gaagtcttca tctgatgtc 180
tgaacaggga aagttttaaag ngagggttga catttcacat ttagcatgtt gtttaacaac 240
ttttcacaag ccgacctga ctttcaggaa gtgaaatgaa aatggcanaa tttatctgaa 300
natccacaat ctaaaaatgg a 321
```

<210> 299

<211> 401

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 104, 268, 347

<223> n = A,T,C or G

<400> 299

```
tatcataaag agtggtgaag tttatttatt atagcaccat tgagacattt tgaaattgga 60
attggtaaaa aaataaaaca aaaagcattt gaattgtatt tggnggaaca gcaaaaaaag 120
agaagtatca tttttctttg tcaaattata ctgtttccaa acattttgga aataaataac 180
tggaattttg tcggctcactt gcaactgggtt acaagattag aacaagagga acacatatgg 240
agttaaattt tttttgttgg gatttcanat agagtttggg ttataaaaag caaacagggc 300
caacgtccac accaaattct tgatcaggac caccaatgtc atagggngca atatctacaa 360
taggtagtct cacagccttg cgtgttcgat attcaaagac t 401
```

<210> 300

<211> 188  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 48  
 <223> n = A,T,C or G

<400> 300  
 tgaatgcttt gtcataattaa gaaagttaaa gtgcaataat gtttgaanac aataagtgg 60  
 ggtgtatctt gtttctaata agataaacctt ttttgtcttt gctttatctt attagggagt 120  
 tgtatgtcag tgtataaaac atactgtgtg gtataacagg cttataaat tctttaaaag 180  
 gaaaaaaa 188

<210> 301  
 <211> 291  
 <212> DNA  
 <213> Homo sapiens

<400> 301  
 aagattttgt tttattttat tatggctaga aagacactgt tatagccaaa atcggcaatg 60  
 acactaaaga aatcctctgt gcttttcaat atgcaaata atttcttcca agagttgccc 120  
 tgggtgtgact tcaagagttc atgttaactt cttttctgga aacttccttt tcttagttgt 180  
 tgtattcttg aagagcctgg gccatgaaga gcttgccaa gttttgggca gtgaactcct 240  
 tgatgttctg gcagtaagtg tttatctggc ctgcaatgag cagcgagtcc a 291

<210> 302  
 <211> 341  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 25  
 <223> n = A,T,C or G

<400> 302  
 tgatttttca taattttatt aaatnatcac tgggaaaact aatgggttcgc gtatcacaca 60  
 attacactac aatctgatag gagtggtaaa accagccaat ggaatccagg taaagtacaa 120  
 aaacgccacc ttttattgtc ctgtcttatt tctcggaag gagggttcta ctttacacat 180  
 ttcattgagcc agcagtggac ttgagttaca atgtgtagg tcttctgtgt tatagctgca 240  
 gaagaagcca tcaaattctt gaggacttga catctctcgg aaagaagcaa actagtggat 300  
 cccccgggct gcaggaattc gatatcaagc ttatcgatac c 341

<210> 303  
 <211> 361  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 15, 27, 92, 124, 127, 183, 198, 244, 320  
 <223> n = A,T,C or G

```

<400> 303
tgcagacagt aaatnaattt tatttgnngtt cacagaacat actaggcgat ctcgacagtc 60
gctccgtgac agcccaccaa cccccaaccc tntacctcgc agccacccta aaggcgactt 120
caanaanatg gaaggatctc acggatctca ttcctaattg tccgccgaag tctcacacag 180
tanacagacg gagttganat gctggaggat gcagtcacct cctaaactta cgacccacca 240
ccanacttca tcccagccgg gacgtcctcc cccacccgag tcctccccat ttcttctcct 300
actttgccgc agttccaggn gtctgtcttc caccagtccc acaaagctca ataaatatca 360
a 361

```

<210> 304

<211> 301

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 23, 104, 192

<223> n = A,T,C or G

<400> 304

```

ctctttacaa cagcctttat ttncggccct tgatcctgct cggatgctgg tggaggccct 60
tagtccgcc cgccaggctc tgtgccgcct ccccgaggc gcanattcat gaacacgggtg 120
ctcaggggct tgaggccgta ctccccagc gggagctggt cctccagggg cttccccctcg 180
aaggtcagcc anaacaggct gtcctgcaca ccctccagcc cgctcacttg ctgcttcagg 240
tgggccacgg tctgcgtcag ccgcacctcg taggtgctgc tgcggccctt gttattcctc 300
a 301

```

<210> 305

<211> 331

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 3, 36, 60, 193, 223

<223> n = A,T,C or G

<400> 305

```

ganaggctag taacatcagt tttattgggt tggggnggca accatagcct ggctgggggn 60
ggggctggcc ctcacagggt gttgagttcc agcagggtct ggtccaaggt ctggtgaatc 120
tcgacgttct ctccttggc actggccaag gtctcttcta ggtcatcgat ggttttctcc 180
aactttgcc canacctctc ggcaaactct gctcgggtct canctcctt cagcttctcc 240
tccaacagtt tgatctcctc ttcataattt tcttctttgg ggaataactc ctctctgag 300
gccatcaggg acttgagggc ctggtccatg g 331

```

<210> 306

<211> 457

<212> DNA

<213> Homo sapiens

<400> 306

```

aatatgtaaa ggtaataact tttattatat taaagacaat gcaaacgaaa aacagaattg 60
agcagtgcaa aatttaaagg actgttttgt tctcaaagtt gcaagtttca aagccaaaag 120

```



```

aattatatgt atcaaatata taagtaaaaa aaagttagac tttcaagcct gtaatcccag 180
cacttttgga ggctgaggca ggtggatcac taacattaaa aagacaacat tagattttgt 240
cgatttatag caattttata aatatataac tttgtcactt ggatcctgaa gcaaaaataat 300
aaagtgaatt tgggattttt gtacttggtg aaaagtttaa caccctaaat tcacaactag 360
tggatccccg gggctgcagg aattcgatat caagcttatc gataccgtcg acctcgaggg 420
ggggcccggg acccaattcg ccctatagtg agtcgta 457

```

<210> 307

<211> 491

<212> DNA

<213> Homo sapiens

<400> 307

```

gtgcttgga ggaacccggc gctcgttccc cccccgggc ggccgcccac agccagccct 60
ccgtcacctc ttcaccgcac cctcggactg cccaaggcc ccgcccggcg ctccagcgcc 120
gcgcagccac cgccgcccgc gccgctctc cttagtgcgc gccatgacga ccgctgccac 180
ctcgcaggtg cgccagaact accaccagga ctcagaggcc gccatcaacc gccagatcaa 240
cctggagctc tacgcctcct acgtttacct gtccatgtct tactactttg accgcgatga 300
tgtggctttg aagaactttg ccaaatactt tcttcaccaa tctcatgagg agagggaaca 360
tgctgagaaa ctgatgaagc tgcagaacca acgagggtggc cgaatcttcc ttcaggatat 420
caagaaacca gactgtgatg actgggagag cgggctgaat gcaatggagt gtgcattaca 480
tttggaaaaa a 491

```

<210> 308

<211> 421

<212> DNA

<213> Homo sapiens

<400> 308

```

ctcagcgctt cttctttctt ggtttgatcc tgactgctgt catggcgctgc cctctggaga 60
aggccctgga tgtgatggtg tccaccttcc acaagtactc gggcaaagag ggtgacaagt 120
tcaagctcaa caagtcagaa ctaaaggagc tgctgacctg ggagctgccc agcttcttgg 180
ggaaaaggac agatgaagct gctttccaga agctgatgag caacttggac agcaacaggg 240
acaacgaggt ggacttccaa gagtactgtg tcttcctgtc ctgcatcgcc atgatgtgta 300
acgaattctt tgaaggcttc ccagataagc agcccaggaa gaaatgaaaa ctcctctgat 360
gtggttgggg ggtctgccag ctggggccct ccctgtcgcc agtgggcact tttttttttc 420
c 421

```

<210> 309

<211> 321

<212> DNA

<213> Homo sapiens

<400> 309

```

accaaaatggc ggatgacgcc ggtgcagcgg gggggcccgg gggccctggt gggcctggga 60
tggggaaccg cggtggcttc cgcggagggt tcggcagtgg catccggggc cggggtcgcg 120
gccgtggacg gggccggggc cgaggccgcg gagctcgcgg aggcaaggcc gaggataagg 180
agtggatgcc cgtcaccaag ttgggcccgt tggtaagga catgaagatc aagtccctgg 240
aggagatcta tctcttctcc ctgccatta aggaatcaga gatcattgat ttcttcctgg 300
ggcctctct caaggatgag g 321

```

<210> 310

<211> 381

<212> DNA

<213> Homo sapiens

<400> 310

```
ttaaccagcc atattggctc aataaatagc ttcggttaagg agttaatttc cttctagaaa 60
tcagtgccta tttttcctgg aaactcaatt ttaaatagtc caattccatc tgaagccaag 120
ctgttgatcat tttcattcgg tgacattctc tcccatgaca cccagaaggg gcagaagaac 180
cacatttttc atttatagat gtttgcattc tttgtattaa aattattttg aaggggttgc 240
ctcattggat ggcttttttt tttttcctcc agggagaagg ggagaaatgt acttggaat 300
taatgtatgt ttacatctct ttgcaaattc ctgtacatag agatatattt tttaagtgtg 360
aatgtaacaa catactgtga a 381
```

<210> 311

<211> 538

<212> DNA

<213> Homo sapiens

<400> 311

```
tttgaattta caccaagaac ttctcaataa aagaaaatca tgaatgctcc acaatttcaa 60
cataccacaa gagaagttaa tttcttaaca ttgtgttcta tgattatttg taagaccttc 120
accaagtctt gatattcttt aaagacatag ttcaaaattg cttttgaaaa tctgtattct 180
tgaaaatatc cttgttggtg attagggttt taaataaccag cttaaaggatt acctcactga 240
gtcatcagta ccttcctatt cagctcccca agatgatgtg tttttgctta cctaagaga 300
ggttttcttc ttatttttag ataattcaag tgcttagata aattatgttt tctttaagtg 360
tttatggtaa actcttttaa agaaaattta atatgttata gctgaatctt tttggtaact 420
ttaaactctt atcatagact ctgtacatat gttcaaatia gctgcttgcc tgatgtgtgt 480
atcatcggtg ggatgacaga acaaacatat ttatgatcat gaataatgtg ctttgtaa 538
```

<210> 312

<211> 176

<212> DNA

<213> Homo sapiens

<400> 312

```
ggaggagcag ctgagagata gggtcagtga atgcggttca gcctgctacc tctcctgtct 60
tcatagaacc attgccttag aattattgta tgacacgttt tttgttggtt aagctgtaag 120
gttttggtct ttgtgaacat gggatattttg aggggagggg ggaggagta gggaag 176
```

<210> 313

<211> 396

<212> DNA

<213> Homo sapiens

<400> 313

```
ccagcaccac caggccctgg gggacctggg ttctcagact gccaaagaag ccttgccatc 60
tggcgctccc atggctcttg caacatctcc ccttcgtttt tgaggggggtc atgccggggg 120
agccaccagc cctcactggg gttcggagga gagtcaggaa gggccaagca cgacaaagca 180
gaaacatcgg atttggggaa cgcgtgtcaa tcccttgtgc cgcagggtcg ggccggagag 240
actgttctgt tcttgtgtga actgtgttgc tgaaagacta cctcgttctt gtcttgatgt 300
gtcaccgggg caactgcctg ggggcgggga tgggggcagg gtggaagcgg ctccccattt 360
tataccaaag gtgctacatc tatgtgatgg gtgggg 396
```

<210> 314

<211> 311

<212> DNA

<213> Homo sapiens

<400> 314

```
cctcaacatc ctcagagagg actggaagcc agtccttacg ataaactcca taatttatgg 60
cctgcagtat ctcttcttgg agcccaaccc cgaggacca ctgaacaagg aggccgcaga 120
ggtcctgcag aacaaccggc ggctgtttga gcagaacgtg cagcgctcca tgcggggtgg 180
ctacatcggc tccacctact ttgagcgctg cctgaaatag ggttggcgca taccaccccc 240
cgccacggcc acaagccctg gcacccctg caaatattta ttgggggcca tgggtagggg 300
tttggggggc g 311
```

<210> 315

<211> 336

<212> DNA

<213> Homo sapiens

<400> 315

```
tttagaacat ggttatcatc caagactact ctaccctgca acattgaact cccaagagca 60
aatccacatt cctcttgagt tctgcagctt ctgtgtaaat agggcagctg tcgtctatgc 120
cgtagaatca catgatctga ggaccattca tggaaagctgc taaatagcct agtctgggga 180
gtcttccata aagttttgca tggagcaaac aaacaggatt aaactagggt tggttccttc 240
agccctctaa aagcataggg cttagcctgc aggcttcctt gggctttctc tgtgtgtgta 300
gttttgtaaa cactatagca tctgttaaga tccagt 336
```

<210> 316

<211> 436

<212> DNA

<213> Homo sapiens

<400> 316

```
aacatgggtc gcgtgcctta agagagacgc ttcctgcaga acaggacctg actacaaaga 60
atgtttccat tggaaattgtt ggtaaagact tggagtttac aatctatgat gatgatgatg 120
tgtctccatt cctggaagggt cttgaagaaa gaccacagag aaaggcacag cctgctcaac 180
ctgctgatga acctgcagaa aaggctgatg aaccaatgga acattaagtg ataagccagt 240
ctatatatgt attatcaaat atgtaagaat acaggcacca catactgatg acaataatct 300
atactttgaa ccaaaagttg cagagtgggtg gaatgctatg ttttaggaat cagtccagat 360
gtgagttttt tccaagcaac ctcaactgaaa cctatataat ggaatacatt tttctttgaa 420
agggctctgta taatca 436
```

<210> 317

<211> 196

<212> DNA

<213> Homo sapiens

<400> 317

```
tattccttgt gaagatgata tactatTTTT gttaagcgtg tctgtattta tgtgtgagga 60
gctgctggct tgcagtgcgc gtgcacgtgg agagctgggtg cccggagatt ggacggcctg 120
atgctccctc cctgcccctg gtccagggaa gctggccgag ggtcctggct cctgaggggc 180
atctgcccct ccccca 196
```

<210> 318

<211> 381

<212> DNA

<213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> 8, 9, 102, 122, 167, 182, 193, 235, 253, 265, 266, 290, 321, 378  
 <223> n = A,T,C or G

<400> 318  
 gacgcttngg ccgtaacgat gatcggagac atcctgctgt tcgggacgtt gctgatgaat 60  
 gccggggcgg tgctgaactt taagctgaaa aagaaggaca cncagggtt tggggaggag 120  
 tncagggagc ccaacacagg tgacaacatc cggaattct tgctgancct cagatacttt 180  
 cnaatcttca tncacctgtg gaacatcttc atgatgttct gcatgattgt gctgntcggc 240  
 tcttgaatcc cancgatgaa accannaact cactttcccg ggatgccgan tctccattcc 300  
 tccattcctg atgacttcaa naatgttttt gacaaaaaaa ccgacaacct tcccagaaag 360  
 tccaagctcg tggtagggngg a 381

<210> 319  
 <211> 506  
 <212> DNA  
 <213> Homo sapiens

<400> 319  
 ctaagcttta cgaatggggt gacaacttat gataaaaact agagctagt g aattagccta 60  
 tttgtaaaata cctttgttat aattgatagg atacatcttg gacatggaat t gtttaagcca 120  
 cctctgagca gtgtatgtca ggacttggtc attaggttgg cagcagaggg g cagaaggaa 180  
 ttatacaggt agagatgtat gcagatgtgt ccatatatgt ccatatttac attttgatag 240  
 ccattgatgt atgcatctct tggctgtact ataagaacac attaatcaa tggaaatata 300  
 ctttgcta attttaattgg tatagatctg ctaatgaatt ctcttaaaaa catactgtat 360  
 tctgttgctg tgtgtttcat tttaaattga gcattaaggg aatgcagcat ttaaatacaga 420  
 actctgccaa tgcttttatc tagaggcgtg ttgccatitt tgtcttatat gaaattttctg 480  
 tccaagaaa ggcaggatta catctt 506

<210> 320  
 <211> 351  
 <212> DNA  
 <213> Homo sapiens

<400> 320  
 ctgacctgca ggacgaaaacc atgaagagcc tgatccttct tgccatcctg gccgccttag 60  
 cggtagtaac tttgtgttat gaatcacatg aaagcatgga atcttatgaa cttaatccct 120  
 tcattaacag gagaaatgca aataccttca tatcccctca gcagagatgg agagctaaag 180  
 tccaagagag gatccgagaa cgctctaagc ctgtccacga gctcaatagg gaagcctgtg 240  
 atgactacag actttgcgaa cgctacgcca tggtttatgg atacaatgct gcctataatc 300  
 gctacttcag gaagcgccga gggaccaaat gagactgagg gaagaaaaaa a 351

<210> 321  
 <211> 421  
 <212> DNA  
 <213> Homo sapiens

<400> 321  
 ctcgaggcgg ttcagctgct tcaagatgaa gctgaacatc tccttcccag ccaactggctg 60  
 ccagaaactc attgaagtgg acgatgaacg caaacttcgt actttctatg agaagcgtat 120  
 ggccacagaa gttgctgctg acgctctggg tgaagaatgg aagggttatg tgggccgaat 180  
 cagtgggtggg aacgacaaac aagggtttccc catgaagcag ggtgtcttga cccatggccc 240

```

tgtccgcctg ctactgagta aggggcattc ctgttacaga ccaaggagaa ctggagaaa 300
aaagagaaaa tcagttcgtg gttgcattgt ggatgcaaat ctgagcgttc tcaacttgg 360
tattgtaaaa aaaggagaga aggatattcc tggactgact gatactacag tgcctcgccg 420
c                                                    421

```

<210> 322

<211> 521

<212> DNA

<213> Homo sapiens

<400> 322

```

agcagctctc ctgccacagc tcctcacccc ctgaaaatgt tcgcctgctc caagtttgtc 60
tccactccct ccttgggtcaa gagcacctca cagctgctga gccgtccgt atctgcagt 120
gtgctgaaac gaccggagat actgacagat gagagcctca gcagcttggc agtctcatgt 180
ccccttacct cacttgtctc tagccgcagc ttccaaacca gcgccatttc aagggaacatc 240
gacacagcag ccaagttcat tggagctggg gctgccacag ttgggggtggc tggttctggg 300
gctgggattg gaactgtgtt tgggagcctc atcattgggtt atgccaggaa cccttctctg 360
aagcaacagc tcttctccta cgccattctg ggctttgccc tctcgagggc catggggctc 420
ttttgtctga tggtagcctt tctcatcctc tttgccatgt gaaggagccg tctccacctc 480
ccatagttct cccgcgtctg gttggccccg tgtgttcctt t                                                    521

```

<210> 323

<211> 435

<212> DNA

<213> Homo sapiens

<400> 323

```

ccgaggtcgc acgcgtgaga cttctccgcc gcagacgccg ccgcgatgcg ctacgtcgcc 60
tcctacctgc tggctgccct agggggcaac tcctccccc gccccaagga catcaagaag 120
atcttggaac gcgtgggtat cgaggcggac gacgaccggc tcaacaaggt tatcagtga 180
ctgaatggaa aaaacattga agacgtcatt gcccaaggta ttggcaagct tgccagtgt 240
cctgctggtg gggctgtagc cgtctctgct gccccaggct ctgcagccc tgcgtctgg 300
tctgccccctg ctgcagcaga ggagaagaaa gatgagaaga aggaggagtc tgaagagtca 360
gatgatgaca tgggatttgg cctttttgat taaattcctg cccccctgca aataaagcct 420
ttttacacat ctcaa                                                    435

```

<210> 324

<211> 521

<212> DNA

<213> Homo sapiens

<400> 324

```

aggagatcga ctttcggtgc ccgcaagacc agggctggaa cgccgagatc acgctgcaga 60
tgggtgcagta caagaatcgt caggccatcc tggcgggtcaa atccacgcgg cagaagcagc 120
agcacctggg ccagcagcag cccccctcgc agccgcagcc gcagccgcag ctccagcccc 180
aaccocagcg tcagcctcag ccgcaacccc agcccaatc acaaccccag cctcagcccc 240
aacccaagcc tcagccccag cagctccacc cgtatccgca tccacatcca catccacact 300
ctcatcctca ctgcaccca caccctcacc cgcaccgcga tccgcaccaa ataccgcacc 360
cacaccaca gccgcactcg cagccgcacg ggcaccggct tctccgcagc acctccaact 420
ctgcctgaaa ggggcagctc ccgggcaaga caagggtttg aggacttgag gaagtgggac 480
gagcacattt ctattgtctt cacttggatc aaaagcaaaa c                                                    521

```

<210> 325

<211> 451

<212> DNA

<213> Homo sapiens

<400> 325

```

atttttcattt ccattaacct ggaagctttc atgaatattc tcttctttta aaacatttta 60
acattatttta aacagaaaaa gatgggctct ttctgggttag ttgttacatg atagcagaga 120
tatttttact tagattactt tgggaatgag agattgttgt cttgaactct ggcaactgtac 180
agtgaatgtg tctgtagttg tgttagtttg cattaagcat gtataacatt caagtatgtc 240
atccaaataa gaggcataata cattgaattg tttttaatcc tctgacaagt tgactcttcg 300
acccccaccc ccaccaaga cattttaata gtaaataagag agagagagaa gagttaatga 360
acatgaggta gtgttccact ggcaggatga cttttcaata gctcaaata atttcagtgc 420
ctttatcact tgaattatta acttaatttg a 451

```

<210> 326

<211> 421

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 296

<223> n = A,T,C or G

<400> 326

```

cgcggtcgta agggctgagg atttttggtc cgcacgctcc tgctcctgac tcaccgctgt 60
tcgctctcgc cgaggaacaa gtcggtcagg aagcccgccg gcaacagcca tggcttttaa 120
ggataccgga aaaacacccg tggagccgga ggtggcaatt caccgaattc gaatcaccc 180
aacaagccgc aacgtaaaat ccttggaaaa ggtgtgtgct gacttgataa gaggcgcaaa 240
agaaaagaat ctcaaagtga aaggaccagt tcgaatgcct accaagactt tgagantcac 300
tacaagaaaa actccttggt gtgaagggtc taagacgtgg gatcggttcc agatgagaat 360
tcacaagcga ctcatcgact tgcacagtc cttctgagatt gtttaagcaga ttacttccat 420
c 421

```

<210> 327

<211> 456

<212> DNA

<213> Homo sapiens

<400> 327

```

atcttgacga ggctgcggtg tctgctgcta ttctccgagc ttccgcaatgc cgcctaagga 60
cgacaagaag aagaaggacg ctggaaagtc ggccaagaaa gacaaagacc cagtgaacaa 120
atccgggggc aaggccaaaa agaagaagtg gtccaaaggc aaagttcggg acaagctcaa 180
taacttagtc ttgtttgaca aagctaccta tgataaactc tgtaagggaag ttcccaacta 240
taaacttata accccagctg tggctctctga gagactgaag attcaggagct ccttggccag 300
ggcagccctt caggagctcc ttagtaaagg acttatcaaa ctgggtttcaa agcacagagc 360
tcaagtaatt tacaccagaa ataccaaggg tggagatgct ccagctgctg gtgaagatgc 420
atgaataggt ccaaccagct gtacatttgg aaaaat 456

```

<210> 328

<211> 471

<212> DNA

<213> Homo sapiens

<400> 328

```

gtggaagtga catcgtcttt aaaccctgcg tggcaatccc tgacgcaccg ccgtgatgcc 60
caggggaagac agggcgacct ggaagtccaa ctacttcctt aagatcatcc aactattgga 120
tgattatccg aaatgtttca ttgtgggagc agacaatgtg ggctccaagc agatgcagca 180
gatccgcatg tcccttcgcg ggaaggctgt ggtgctgatg ggcaagaaca ccatgatgcg 240
caaggccatc cgagggcacc tggaaaacaa cccagctctg gagaaactgc tgcctcatat 300
ccgggggaat gtgggctttg ttttcaccaa ggaggacctc actgagatca gggacatgtt 360
gctggccaat aagggtgccag ctgctgcccg tgctggtgcc attgccccat gtgaagtcac 420
tgtgccagcc cagaacactg gtctcgggcc cgagaagacc tcctttttcc a 471

```

<210> 329

<211> 278

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> 154, 204

<223> n = A,T,C or G

<400> 329

```

gtttaaacctt aagcttggtg ccgagctcgg atccactagt ccagtgtggt ggaattctag 60
aaattgagat gcccccccag gccagcaaat gtcccttttt gttcaaagtc tttttttatt 120
ccttgataatt tttctttttt tttttttttt ttgnggatgg ggacttgtga atttttctaa 180
aggtgctatt taacatggga gganagcgtg tgcggctcca gccagccccg ctgctcactt 240
tccacctctc ctccacctgc ctctggtctc tcaggcct 278

```

<210> 330

<211> 338

<212> DNA

<213> Homo sapiens

<400> 330

```

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<210> 331

<211> 2820

<212> DNA

<213> Homo sapiens

<400> 331

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<210> 332

<211> 2270

<212> DNA

<213> Homo sapiens

<400> 332

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<210> 333

<211> 2816

<212> DNA

<213> Homo sapiens

<400> 333

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<210> 334

<211> 2082

<212> DNA

<213> Homo sapiens

<400> 334

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| cactgtagac  | cagggatagc | aaataggcct  | tactataata  | taaagtgact | tgtttgaaatg | 1980 |
| ctgtaatgag  | aagaattctg | agacctagt   | catgataatt  | ggggaaatat | ctgggtgcag  | 2040 |
| aaggataagg  | tagcatcatg | ttgccgtatt  | ttagcatctc  | tg         |             | 2082 |

&lt;210&gt; 335

&lt;211&gt; 4849

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 335

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<211> 1386  
 <212> DNA  
 <213> Homo sapiens

<400> 336

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aacacagacc acgcgcagaa cagcgtcacg gcgcctcgc cctacgcaca gccagctcc 180
accttcgatg ctctctctcc atcaccgcc atccctcca acaccgacta cccaggcccg 240
cacagtttcg acgtgtcctt ccagcagtcg agcaccgcca agtcggccac ctggacgtat 300
tccactgaac tgaagaaact ctactgcaa attgcaaaga catgccccat ccagatcaag 360
gtgatgaccc cacctcctca gggagctgtt atccgcgcca tgctgtcta caaaaaagct 420
gagcacgtca cggagggtgt gaagcgggtg cccaaccatg agctgagccg tgaattcaac 480
gagggacaga ttgccccctc tagtcatttg attcgagtag aggggaacag ccatgcccag 540
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gttggcactg aattcacgac agtcttgtag aatttcatgt gtaacagcag ttgtgttgga 660
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gatgaagata gcatcagaaa gcagcaagtt tcggacagta caaagaacgg tgatggtacg 840
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<210> 337  
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 <212> DNA  
 <213> Homo sapiens

<400> 337

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ccatcagaag atggtgcgac aaacaagatt gagattagca tggactgtat ccgcatgcag 180
gactcggacc tgagtgaccc catgtggcca cagtacacga acctggggct cctgaacagc 240
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cagaacagcg tcacggcgcc ctgcctctac gcacagccca gctccacctt cgatgctctc 360
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aaactctact gccaaattgc aaagacatgc cccatccaga tcaaggatgat gacccacact 540
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gtggtgaagc ggtgccccaa ccatgagctg agccgtgaat tcaacgaggg acagattgcc 660
cctcctagtc atttgattcg agtagagggg aacagccatg cccagtatgt agaagatccc 720
atcacaggaa gacagagtgt gctggtacct tatgagccac cccaggttgg cactgaattc 780
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<210> 338

<211> 586

<212> PRT

<213> Homo sapiens

<400> 338

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Tyr Thr Asn Leu Gly Leu Leu Asn Ser Met Asp Gln Gln Ile Arg Asn
          20          25          30
Gly Ser Ser Ser Thr Ser Pro Tyr Asn Thr Asp His Ala Gln Asn Ser
          35          40          45
Val Thr Ala Pro Ser Pro Tyr Ala Gln Pro Ser Pro Thr Phe Asp Ala
          50          55          60
Leu Ser Pro Ser Pro Ala Ile Pro Ser Asn Thr Asp Tyr Pro Gly Pro
          65          70          75          80
His Ser Ser Asp Val Ser Phe Gln Gln Ser Ser Thr Ala Lys Ser Ala
          85          90          95
Thr Trp Thr Tyr Ser Thr Glu Leu Lys Lys Leu Tyr Cys Gln Ile Ala
          100         105         110
Lys Thr Cys Pro Ile Gln Ile Lys Val Met Thr Pro Pro Pro Gln Gly
          115         120         125
Ala Val Ile Arg Ala Met Pro Val Tyr Lys Lys Ala Glu His Val Thr
          130         135         140
Glu Val Val Lys Arg Cys Pro Asn His Glu Leu Ser Arg Glu Phe Asn
          145         150         155         160
Glu Gly Gln Ile Ala Pro Pro Ser His Leu Ile Arg Val Glu Gly Asn
          165         170         175
Ser His Ala Gln Tyr Val Glu Asp Pro Ile Thr Gly Arg Gln Ser Val
          180         185         190
Leu Val Pro Tyr Glu Pro Pro Gln Val Gly Thr Glu Phe Thr Thr Val
          195         200         205
Leu Tyr Asn Phe Met Cys Asn Ser Ser Cys Val Gly Gly Met Asn Arg
          210         215         220
Arg Pro Ile Leu Ile Ile Val Thr Leu Glu Thr Arg Asp Gly Gln Val
          225         230         235         240
Leu Gly Arg Arg Cys Phe Glu Ala Arg Ile Cys Ala Cys Pro Gly Arg
          245         250         255
Asp Arg Lys Ala Asp Glu Asp Ser Ile Arg Lys Gln Gln Val Ser Asp
          260         265         270
Ser Thr Lys Asn Gly Asp Gly Thr Lys Arg Pro Phe Arg Gln Asn Thr
          275         280         285
His Gly Ile Gln Met Thr Ser Ile Lys Lys Arg Arg Ser Pro Asp Asp
          290         295         300
Glu Leu Leu Tyr Leu Pro Val Arg Gly Arg Glu Thr Tyr Glu Met Leu

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305          310          315          320
Leu Lys Ile Lys Glu Ser Leu Glu Leu Met Gln Tyr Leu Pro Gln His
          325          330          335
Thr Ile Glu Thr Tyr Arg Gln Gln Gln Gln Gln Gln His Gln His Leu
          340          345          350
Leu Gln Lys Gln Thr Ser Ile Gln Ser Pro Ser Ser Tyr Gly Asn Ser
          355          360          365
Ser Pro Pro Leu Asn Lys Met Asn Ser Met Asn Lys Leu Pro Ser Val
          370          375          380
Ser Gln Leu Ile Asn Pro Gln Gln Arg Asn Ala Leu Thr Pro Thr Thr
385          390          395          400
Ile Pro Asp Gly Met Gly Ala Asn Ile Pro Met Met Gly Thr His Met
          405          410          415
Pro Met Ala Gly Asp Met Asn Gly Leu Ser Pro Thr Gln Ala Leu Pro
          420          425          430
Pro Pro Leu Ser Met Pro Ser Thr Ser His Cys Thr Pro Pro Pro Pro
          435          440          445
Tyr Pro Thr Asp Cys Ser Ile Val Ser Phe Leu Ala Arg Leu Gly Cys
          450          455          460
Ser Ser Cys Leu Asp Tyr Phe Thr Thr Gln Gly Leu Thr Thr Ile Tyr
465          470          475          480
Gln Ile Glu His Tyr Ser Met Asp Asp Leu Ala Ser Leu Lys Ile Pro
          485          490          495
Glu Gln Phe Arg His Ala Ile Trp Lys Gly Ile Leu Asp His Arg Gln
          500          505          510
Leu His Glu Phe Ser Ser Pro Ser His Leu Leu Arg Thr Pro Ser Ser
          515          520          525
Ala Ser Thr Val Ser Val Gly Ser Ser Glu Thr Arg Gly Glu Arg Val
          530          535          540
Ile Asp Ala Val Arg Phe Thr Leu Arg Gln Thr Ile Ser Phe Pro Pro
545          550          555          560
Arg Asp Glu Trp Asn Asp Phe Asn Phe Asp Met Asp Ala Arg Arg Asn
          565          570          575
Lys Gln Gln Arg Ile Lys Glu Glu Gly Glu
          580          585

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<210> 339
<211> 641
<212> PRT
<213> Homo sapiens

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<400> 339
Met Ser Gln Ser Thr Gln Thr Asn Glu Phe Leu Ser Pro Glu Val Phe
1          5          10          15
Gln His Ile Trp Asp Phe Leu Glu Gln Pro Ile Cys Ser Val Gln Pro
          20          25          30
Ile Asp Leu Asn Phe Val Asp Glu Pro Ser Glu Asp Gly Ala Thr Asn
          35          40          45
Lys Ile Glu Ile Ser Met Asp Cys Ile Arg Met Gln Asp Ser Asp Leu
          50          55          60
Ser Asp Pro Met Trp Pro Gln Tyr Thr Asn Leu Gly Leu Leu Asn Ser
65          70          75          80
Met Asp Gln Gln Ile Gln Asn Gly Ser Ser Ser Thr Ser Pro Tyr Asn

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|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
|     |     |     |     | 85  |     |     |     |     | 90  |     |     |     |     | 95  |     |     |
| Thr | Asp | His |     | Ala | Gln | Asn | Ser | Val | Thr | Ala | Pro | Ser | Pro | Tyr | Ala | Gln |
|     |     |     |     | 100 |     |     |     |     | 105 |     |     |     |     | 110 |     |     |
| Pro | Ser | Ser | Thr | Phe | Asp | Ala | Leu | Ser | Pro | Ser | Pro | Ala | Ile | Pro | Ser |     |
|     |     |     | 115 |     |     |     | 120 |     |     |     |     | 125 |     |     |     |     |
| Asn | Thr | Asp | Tyr | Pro | Gly | Pro | His | Ser | Phe | Asp | Val | Ser | Phe | Gln | Gln |     |
|     | 130 |     |     |     | 135 |     |     |     |     | 140 |     |     |     |     |     |     |
| Ser | Ser | Thr | Ala | Lys | Ser | Ala | Thr | Trp | Thr | Tyr | Ser | Thr | Glu | Leu | Lys |     |
| 145 |     |     |     | 150 |     |     |     |     |     | 155 |     |     |     |     | 160 |     |
| Lys | Leu | Tyr | Cys | Gln | Ile | Ala | Lys | Thr | Cys | Pro | Ile | Gln | Ile | Lys | Val |     |
|     |     |     | 165 |     |     |     |     |     | 170 |     |     |     |     | 175 |     |     |
| Met | Thr | Pro | Pro | Pro | Gln | Gly | Ala | Val | Ile | Arg | Ala | Met | Pro | Val | Tyr |     |
|     |     |     | 180 |     |     |     |     | 185 |     |     |     |     | 190 |     |     |     |
| Lys | Lys | Ala | Glu | His | Val | Thr | Glu | Val | Val | Lys | Arg | Cys | Pro | Asn | His |     |
|     |     | 195 |     |     |     |     | 200 |     |     |     |     | 205 |     |     |     |     |
| Glu | Leu | Ser | Arg | Glu | Phe | Asn | Glu | Gly | Gln | Ile | Ala | Pro | Pro | Ser | His |     |
|     | 210 |     |     |     | 215 |     |     |     |     |     | 220 |     |     |     |     |     |
| Leu | Ile | Arg | Val | Glu | Gly | Asn | Ser | His | Ala | Gln | Tyr | Val | Glu | Asp | Pro |     |
| 225 |     |     |     | 230 |     |     |     |     |     | 235 |     |     |     |     | 240 |     |
| Ile | Thr | Gly | Arg | Gln | Ser | Val | Leu | Val | Pro | Tyr | Glu | Pro | Pro | Gln | Val |     |
|     |     |     | 245 |     |     |     |     |     | 250 |     |     |     |     | 255 |     |     |
| Gly | Thr | Glu | Phe | Thr | Thr | Val | Leu | Tyr | Asn | Phe | Met | Cys | Asn | Ser | Ser |     |
|     |     |     | 260 |     |     |     |     | 265 |     |     |     |     |     | 270 |     |     |
| Cys | Val | Gly | Gly | Met | Asn | Arg | Arg | Pro | Ile | Leu | Ile | Ile | Val | Thr | Leu |     |
|     |     | 275 |     |     |     | 280 |     |     |     |     |     | 285 |     |     |     |     |
| Glu | Thr | Arg | Asp | Gly | Gln | Val | Leu | Gly | Arg | Arg | Cys | Phe | Glu | Ala | Arg |     |
|     | 290 |     |     |     | 295 |     |     |     |     |     | 300 |     |     |     |     |     |
| Ile | Cys | Ala | Cys | Pro | Gly | Arg | Asp | Arg | Lys | Ala | Asp | Glu | Asp | Ser | Ile |     |
| 305 |     |     |     | 310 |     |     |     |     |     | 315 |     |     |     |     | 320 |     |
| Arg | Lys | Gln | Gln | Val | Ser | Asp | Ser | Thr | Lys | Asn | Gly | Asp | Gly | Thr | Lys |     |
|     |     |     | 325 |     |     |     |     |     | 330 |     |     |     |     | 335 |     |     |
| Arg | Pro | Phe | Arg | Gln | Asn | Thr | His | Gly | Ile | Gln | Met | Thr | Ser | Ile | Lys |     |
|     |     | 340 |     |     |     |     |     | 345 |     |     |     |     | 350 |     |     |     |
| Lys | Arg | Arg | Ser | Pro | Asp | Asp | Glu | Leu | Leu | Tyr | Leu | Pro | Val | Arg | Gly |     |
|     |     | 355 |     |     |     |     | 360 |     |     |     |     | 365 |     |     |     |     |
| Arg | Glu | Thr | Tyr | Glu | Met | Leu | Leu | Lys | Ile | Lys | Glu | Ser | Leu | Glu | Leu |     |
|     | 370 |     |     |     | 375 |     |     |     |     |     | 380 |     |     |     |     |     |
| Met | Gln | Tyr | Leu | Pro | Gln | His | Thr | Ile | Glu | Thr | Tyr | Arg | Gln | Gln | Gln |     |
| 385 |     |     |     | 390 |     |     |     |     |     | 395 |     |     |     |     | 400 |     |
| Gln | Gln | Gln | His | Gln | His | Leu | Leu | Gln | Lys | Gln | Thr | Ser | Ile | Gln | Ser |     |
|     |     |     | 405 |     |     |     |     |     | 410 |     |     |     |     | 415 |     |     |
| Pro | Ser | Ser | Tyr | Gly | Asn | Ser | Ser | Pro | Pro | Leu | Asn | Lys | Met | Asn | Ser |     |
|     |     | 420 |     |     |     |     |     | 425 |     |     |     |     | 430 |     |     |     |
| Met |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |



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<210> 340
<211> 448
<212> PRT
<213> Homo sapiens
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|          |     |     |     |          |     |     |     |     |           |     |     |     |     |           |     |
|----------|-----|-----|-----|----------|-----|-----|-----|-----|-----------|-----|-----|-----|-----|-----------|-----|
| Met<br>1 | Ser | Gln | Ser | Thr<br>5 | Gln | Thr | Asn | Glu | Phe<br>10 | Leu | Ser | Pro | Glu | Val<br>15 | Phe |
| Gln      | His | Ile | Trp | Asp      | Phe | Leu | Glu | Gln | Pro       | Ile | Cys | Ser | Val | Gln       | Pro |
|          |     |     | 20  |          |     |     |     | 25  |           |     |     |     | 30  |           |     |
| Ile      | Asp | Leu | Asn | Phe      | Val | Asp | Glu | Pro | Ser       | Glu | Asp | Gly | Ala | Thr       | Asn |
|          |     |     | 35  |          |     |     | 40  |     |           |     |     | 45  |     |           |     |
| Lys      | Ile | Glu | Ile | Ser      | Met | Asp | Cys | Ile | Arg       | Met | Gln | Asp | Ser | Asp       | Leu |
|          |     |     | 50  |          |     | 55  |     |     |           |     | 60  |     |     |           |     |
| Ser      | Asp | Pro | Met | Trp      | Pro | Gln | Tyr | Thr | Asn       | Leu | Gly | Leu | Leu | Asn       | Ser |
| 65       |     |     |     |          | 70  |     |     |     |           | 75  |     |     |     |           | 80  |
| Met      | Asp | Gln | Gln | Ile      | Gln | Asn | Gly | Ser | Ser       | Ser | Thr | Ser | Pro | Tyr       | Asn |
|          |     |     |     | 85       |     |     |     | 90  |           |     |     |     |     | 95        |     |
| Thr      | Asp | His | Ala | Gln      | Asn | Ser | Val | Thr | Ala       | Pro | Ser | Pro | Tyr | Ala       | Gln |
|          |     |     | 100 |          |     |     |     | 105 |           |     |     |     | 110 |           |     |
| Pro      | Ser | Ser | Thr | Phe      | Asp | Ala | Leu | Ser | Pro       | Ser | Pro | Ala | Ile | Pro       | Ser |
|          |     |     | 115 |          |     |     | 120 |     |           |     |     | 125 |     |           |     |
| Asn      | Thr | Asp | Tyr | Pro      | Gly | Pro | His | Ser | Phe       | Asp | Val | Ser | Phe | Gln       | Gln |
|          |     |     |     |          | 130 |     | 135 |     |           |     | 140 |     |     |           |     |
| Ser      | Ser | Thr | Ala | Lys      | Ser | Ala | Thr | Trp | Thr       | Tyr | Ser | Thr | Glu | Leu       | Lys |
| 145      |     |     |     |          | 150 |     |     |     |           | 155 |     |     |     |           | 160 |
| Lys      | Leu | Tyr | Cys | Gln      | Ile | Ala | Lys | Thr | Cys       | Pro | Ile | Gln | Ile | Lys       | Val |
|          |     |     |     | 165      |     |     |     |     | 170       |     |     |     |     | 175       |     |
| Met      | Thr | Pro | Pro | Pro      | Gln | Gly | Ala | Val | Ile       | Arg | Ala | Met | Pro | Val       | Tyr |
|          |     |     | 180 |          |     |     |     | 185 |           |     |     |     | 190 |           |     |
| Lys      | Lys | Ala | Glu | His      | Val | Thr | Glu | Val | Val       | Lys | Arg | Cys | Pro | Asn       | His |
|          |     | 195 |     |          |     |     | 200 |     |           |     |     | 205 |     |           |     |
| Glu      | Leu | Ser | Arg | Glu      | Phe | Asn | Glu | Gly | Gln       | Ile | Ala | Pro | Pro | Ser       | His |
|          |     |     |     |          |     | 215 |     |     |           |     | 220 |     |     |           |     |
| Leu      | Ile | Arg | Val | Glu      | Gly | Asn | Ser | His | Ala       | Gln | Tyr | Val | Glu | Asp       | Pro |

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225          230          235          240
Ile Thr Gly Arg Gln Ser Val Leu Val Pro Tyr Glu Pro Pro Gln Val
          245          250          255
Gly Thr Glu Phe Thr Thr Val Leu Tyr Asn Phe Met Cys Asn Ser Ser
          260          265          270
Cys Val Gly Gly Met Asn Arg Arg Pro Ile Leu Ile Ile Val Thr Leu
          275          280          285
Glu Thr Arg Asp Gly Gln Val Leu Gly Arg Arg Cys Phe Glu Ala Arg
          290          295          300
Ile Cys Ala Cys Pro Gly Arg Asp Arg Lys Ala Asp Glu Asp Ser Ile
305          310          315          320
Arg Lys Gln Gln Val Ser Asp Ser Thr Lys Asn Gly Asp Gly Thr Lys
          325          330          335
Arg Pro Phe Arg Gln Asn Thr His Gly Ile Gln Met Thr Ser Ile Lys
          340          345          350
Lys Arg Arg Ser Pro Asp Asp Glu Leu Leu Tyr Leu Pro Val Arg Gly
          355          360          365
Arg Glu Thr Tyr Glu Met Leu Leu Lys Ile Lys Glu Ser Leu Glu Leu
          370          375          380
Met Gln Tyr Leu Pro Gln His Thr Ile Glu Thr Tyr Arg Gln Gln Gln
385          390          395          400
Gln Gln Gln His Gln His Leu Leu Gln Lys His Leu Leu Ser Ala Cys
          405          410          415
Phe Arg Asn Glu Leu Val Glu Pro Arg Arg Glu Thr Pro Lys Gln Ser
          420          425          430
Asp Val Phe Phe Arg His Ser Lys Pro Pro Asn Arg Ser Val Tyr Pro
          435          440          445

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<210> 341

<211> 356

<212> PRT

<213> Homo sapiens

<400> 341

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Met Leu Tyr Leu Glu Asn Asn Ala Gln Thr Gln Phe Ser Glu Pro Gln
 1          5          10          15
Tyr Thr Asn Leu Gly Leu Leu Asn Ser Met Asp Gln Gln Ile Gln Asn
          20          25          30
Gly Ser Ser Ser Thr Ser Pro Tyr Asn Thr Asp His Ala Gln Asn Ser
          35          40          45
Val Thr Ala Pro Ser Pro Tyr Ala Gln Pro Ser Ser Thr Phe Asp Ala
          50          55          60
Leu Ser Pro Ser Pro Ala Ile Pro Ser Asn Thr Asp Tyr Pro Gly Pro
65          70          75          80
His Ser Phe Asp Val Ser Phe Gln Gln Ser Ser Thr Ala Lys Ser Ala
          85          90          95
Thr Trp Thr Tyr Ser Thr Glu Leu Lys Lys Leu Tyr Cys Gln Ile Ala
          100          105          110
Lys Thr Cys Pro Ile Gln Ile Lys Val Met Thr Pro Pro Pro Gln Gly
          115          120          125
Ala Val Ile Arg Ala Met Pro Val Tyr Lys Lys Ala Glu His Val Thr
          130          135          140
Glu Val Val Lys Arg Cys Pro Asn His Glu Leu Ser Arg Glu Phe Asn

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145                      150                      155                      160  
 Glu Gly Gln Ile Ala Pro Pro Ser His Leu Ile Arg Val Glu Gly Asn  
                                  165                      170                      175  
 Ser His Ala Gln Tyr Val Glu Asp Pro Ile Thr Gly Arg Gln Ser Val  
                                  180                      185                      190  
 Leu Val Pro Tyr Glu Pro Pro Gln Val Gly Thr Glu Phe Thr Thr Val  
                                  195                      200                      205  
 Leu Tyr Asn Phe Met Cys Asn Ser Ser Cys Val Gly Gly Met Asn Arg  
                                  210                      215                      220  
 Arg Pro Ile Leu Ile Ile Val Thr Leu Glu Thr Arg Asp Gly Gln Val  
 225                      230                      235                      240  
 Leu Gly Arg Arg Cys Phe Glu Ala Arg Ile Cys Ala Cys Pro Gly Arg  
                                  245                      250                      255  
 Asp Arg Lys Ala Asp Glu Asp Ser Ile Arg Lys Gln Gln Val Ser Asp  
                                  260                      265                      270  
 Ser Thr Lys Asn Gly Asp Gly Thr Lys Arg Pro Ser Arg Gln Asn Thr  
                                  275                      280                      285  
 His Gly Ile Gln Met Thr Ser Ile Lys Lys Arg Arg Ser Pro Asp Asp  
                                  290                      295                      300  
 Glu Leu Leu Tyr Leu Pro Val Arg Gly Arg Glu Thr Tyr Glu Met Leu  
 305                      310                      315                      320  
 Leu Lys Ile Lys Glu Ser Leu Glu Leu Met Gln Tyr Leu Pro Gln His  
                                  325                      330                      335  
 Thr Ile Glu Thr Tyr Arg Gln Gln Gln Gln Gln Gln His Gln His Leu  
                                  340                      345                      350  
 Leu Gln Lys Gln  
                                  355

<210> 342  
 <211> 680  
 <212> PRT  
 <213> Homo sapiens

<400> 342  
 Met Asn Phe Glu Thr Ser Arg Cys Ala Thr Leu Gln Tyr Cys Pro Asp  
 1                      5                      10                      15  
 Pro Tyr Ile Gln Arg Phe Val Glu Thr Pro Ala His Phe Ser Trp Lys  
                                  20                      25                      30  
 Glu Ser Tyr Tyr Arg Ser Thr Met Ser Gln Ser Thr Gln Thr Asn Glu  
                                  35                      40                      45  
 Phe Leu Ser Pro Glu Val Phe Gln His Ile Trp Asp Phe Leu Glu Gln  
 50                      55                      60  
 Pro Ile Cys Ser Val Gln Pro Ile Asp Leu Asn Phe Val Asp Glu Pro  
 65                      70                      75                      80  
 Ser Glu Asp Gly Ala Thr Asn Lys Ile Glu Ile Ser Met Asp Cys Ile  
                                  85                      90                      95  
 Arg Met Gln Asp Ser Asp Leu Ser Asp Pro Met Trp Pro Gln Tyr Thr  
                                  100                      105                      110  
 Asn Leu Gly Leu Leu Asn Ser Met Asp Gln Gln Ile Gln Asn Gly Ser  
                                  115                      120                      125  
 Ser Ser Thr Ser Pro Tyr Asn Thr Asp His Ala Gln Asn Ser Val Thr  
 130                      135                      140  
 Ala Pro Ser Pro Tyr Ala Gln Pro Ser Ser Thr Phe Asp Ala Leu Ser

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |  |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|
| 145 |     |     |     |     | 150 |     |     |     | 155 |     |     |     | 160 |     |     |  |
| Pro | Ser | Pro | Ala | Ile | Pro | Ser | Asn | Thr | Asp | Tyr | Pro | Gly | Pro | His | Ser |  |
|     |     |     |     | 165 |     |     |     |     | 170 |     |     |     |     | 175 |     |  |
| Phe | Asp | Val | Ser | Phe | Gln | Gln | Ser | Ser | Thr | Ala | Lys | Ser | Ala | Thr | Trp |  |
|     |     |     | 180 |     |     |     |     | 185 |     |     |     |     | 190 |     |     |  |
| Thr | Tyr | Ser | Thr | Glu | Leu | Lys | Lys | Leu | Tyr | Cys | Gln | Ile | Ala | Lys | Thr |  |
|     |     | 195 |     |     |     |     | 200 |     |     |     |     | 205 |     |     |     |  |
| Cys | Pro | Ile | Gln | Ile | Lys | Val | Met | Thr | Pro | Pro | Pro | Gln | Gly | Ala | Val |  |
|     | 210 |     |     |     | 215 |     |     |     |     | 220 |     |     |     |     |     |  |
| Ile | Arg | Ala | Met | Pro | Val | Tyr | Lys | Lys | Ala | Glu | His | Val | Thr | Glu | Val |  |
| 225 |     |     |     | 230 |     |     |     |     | 235 |     |     |     |     | 240 |     |  |
| Val | Lys | Arg | Cys | Pro | Asn | His | Glu | Leu | Ser | Arg | Glu | Phe | Asn | Glu | Gly |  |
|     |     |     |     | 245 |     |     |     |     | 250 |     |     |     |     | 255 |     |  |
| Gln | Ile | Ala | Pro | Pro | Ser | His | Leu | Ile | Arg | Val | Glu | Gly | Asn | Ser | His |  |
|     |     |     | 260 |     |     |     |     | 265 |     |     |     |     | 270 |     |     |  |
| Ala | Gln | Tyr | Val | Glu | Asp | Pro | Ile | Thr | Gly | Arg | Gln | Ser | Val | Leu | Val |  |
|     |     | 275 |     |     |     |     | 280 |     |     |     |     | 285 |     |     |     |  |
| Pro | Tyr | Glu | Pro | Pro | Gln | Val | Gly | Thr | Glu | Phe | Thr | Thr | Val | Leu | Tyr |  |
|     | 290 |     |     |     | 295 |     |     |     |     |     | 300 |     |     |     |     |  |
| Asn | Phe | Met | Cys | Asn | Ser | Ser | Cys | Val | Gly | Gly | Met | Asn | Arg | Arg | Pro |  |
| 305 |     |     |     |     | 310 |     |     |     |     | 315 |     |     |     |     | 320 |  |
| Ile | Leu | Ile | Ile | Val | Thr | Leu | Glu | Thr | Arg | Asp | Gly | Gln | Val | Leu | Gly |  |
|     |     |     |     | 325 |     |     |     |     | 330 |     |     |     |     | 335 |     |  |
| Arg | Arg | Cys | Phe | Glu | Ala | Arg | Ile | Cys | Ala | Cys | Pro | Gly | Arg | Asp | Arg |  |
|     |     |     | 340 |     |     |     |     | 345 |     |     |     |     | 350 |     |     |  |
| Lys | Ala | Asp | Glu | Asp | Ser | Ile | Arg | Lys | Gln | Gln | Val | Ser | Asp | Ser | Thr |  |
|     |     | 355 |     |     |     |     | 360 |     |     |     |     | 365 |     |     |     |  |
| Lys | Asn | Gly | Asp | Gly | Thr | Lys | Arg | Pro | Phe | Arg | Gln | Asn | Thr | His | Gly |  |
|     | 370 |     |     |     |     | 375 |     |     |     |     | 380 |     |     |     |     |  |
| Ile | Gln | Met | Thr | Ser | Ile | Lys | Lys | Arg | Arg | Ser | Pro | Asp | Asp | Glu | Leu |  |
| 385 |     |     |     |     | 390 |     |     |     |     | 395 |     |     |     | 400 |     |  |
| Leu | Tyr | Leu | Pro | Val | Arg | Gly | Arg | Glu | Thr | Tyr | Glu | Met | Leu | Leu | Lys |  |
|     |     |     |     | 405 |     |     |     |     | 410 |     |     |     |     | 415 |     |  |
| Ile | Lys | Glu | Ser | Leu | Glu | Leu | Met | Gln | Tyr | Leu | Pro | Gln | His | Thr | Ile |  |
|     |     |     | 420 |     |     |     |     | 425 |     |     |     |     | 430 |     |     |  |
| Glu | Thr | Tyr | Arg | Gln | Gln | Gln | Gln | Gln | Gln | His | Gln | His | Leu | Leu | Gln |  |
|     |     | 435 |     |     |     |     |     | 440 |     |     |     | 445 |     |     |     |  |
| Lys | Gln | Thr | Ser | Ile | Gln | Ser | Pro | Ser | Ser | Tyr | Gly | Asn | Ser | Ser | Pro |  |
|     | 450 |     |     |     |     | 455 |     |     |     |     | 460 |     |     |     |     |  |
| Pro | Leu | Asn | Lys | Met | Asn | Ser | Met | Asn | Lys | Leu | Pro | Ser | Val | Ser | Gln |  |
| 465 |     |     |     |     | 470 |     |     |     |     | 475 |     |     |     |     | 480 |  |
| Leu | Ile | Asn | Pro | Gln | Gln | Arg | Asn | Ala | Leu | Thr | Pro | Thr | Thr | Ile | Pro |  |
|     |     |     |     | 485 |     |     |     |     | 490 |     |     |     |     | 495 |     |  |
| Asp | Gly | Met | Gly | Ala | Asn | Ile | Pro | Met | Met | Gly | Thr | His | Met | Pro | Met |  |
|     |     |     | 500 |     |     |     |     | 505 |     |     |     |     | 510 |     |     |  |

<210> 343

&lt;212&gt; PRT

<213> Homo sapiens

<400> 343

|            |     |            |            |          |            |            |            |            |            |            |            |            |            |           |     |
|------------|-----|------------|------------|----------|------------|------------|------------|------------|------------|------------|------------|------------|------------|-----------|-----|
| Met<br>1   | Leu | Tyr        | Leu        | Glu<br>5 | Asn        | Asn        | Ala        | Gln        | Thr<br>10  | Gln        | Phe        | Ser        | Glu        | Pro<br>15 | Gln |
| Tyr        | Thr | Asn        | Leu<br>20  | Gly      | Leu        | Leu        | Asn        | Ser<br>25  | Met        | Asp        | Gln        | Gln        | Ile<br>30  | Gln       | Asn |
| Gly        | Ser | Ser<br>35  | Ser        | Thr      | Ser        | Pro        | Tyr<br>40  | Asn        | Thr        | Asp        | His        | Ala<br>45  | Gln        | Asn       | Ser |
| Val<br>50  | Thr | Ala        | Pro        | Ser      | Pro        | Tyr<br>55  | Ala        | Gln        | Pro        | Ser        | Ser        | Thr<br>60  | Phe        | Asp       | Ala |
| Leu<br>65  | Ser | Pro        | Ser        | Pro      | Ala<br>70  | Ile        | Pro        | Ser        | Asn        | Thr<br>75  | Asp        | Tyr        | Pro        | Gly<br>80 | Pro |
| His        | Ser | Phe        | Asp<br>85  | Val      | Ser        | Phe        | Gln        | Gln        | Ser<br>90  | Ser        | Thr        | Ala        | Lys<br>95  | Ser       | Ala |
| Thr        | Trp | Thr        | Tyr<br>100 | Ser      | Thr        | Glu        | Leu        | Lys<br>105 | Lys        | Leu        | Tyr        | Cys        | Gln<br>110 | Ile       | Ala |
| Lys        | Thr | Cys<br>115 | Pro        | Ile      | Gln        | Ile        | Lys<br>120 | Val        | Met        | Thr        | Pro        | Pro<br>125 | Pro        | Gln       | Gly |
| Ala<br>130 | Val | Ile        | Arg        | Ala      | Met<br>135 | Pro        | Val        | Tyr        | Lys        | Lys        | Ala<br>140 | Glu        | His        | Val       | Thr |
| Glu<br>145 | Val | Val        | Lys        | Arg      | Cys<br>150 | Pro        | Asn        | His        | Glu        | Leu        | Ser<br>155 | Arg        | Glu        | Phe       | Asn |
| Glu        | Gly | Gln        | Ile<br>165 | Ala      | Pro        | Pro        | Ser        | His        | Leu<br>170 | Ile        | Arg        | Val        | Glu<br>175 | Gly       | Asn |
| Ser        | His | Ala        | Gln<br>180 | Tyr      | Val        | Glu        | Asp<br>185 | Pro        | Ile        | Thr        | Gly        | Arg<br>190 | Gln        | Ser       | Val |
| Leu        | Val | Pro<br>195 | Tyr        | Glu      | Pro        | Pro        | Gln<br>200 | Val        | Gly        | Thr        | Glu        | Phe<br>205 | Thr        | Thr       | Val |
| Leu<br>210 | Tyr | Asn        | Phe        | Met      | Cys        | Asn<br>215 | Ser        | Ser        | Cys        | Val        | Gly<br>220 | Gly        | Met        | Asn       | Arg |
| Arg<br>225 | Pro | Ile        | Leu        | Ile      | Ile<br>230 | Val        | Thr        | Leu        | Glu        | Thr<br>235 | Arg        | Asp        | Gly        | Gln       | Val |
| Leu        | Gly | Arg        | Arg<br>245 | Cys      | Phe        | Glu        | Ala        | Arg<br>250 | Ile        | Cys        | Ala        | Cys<br>255 | Pro        | Gly       | Arg |
| Asp        | Arg | Lys        | Ala        | Asp      | Glu        | Asp        | Ser        | Ile        | Arg        | Lys        | Gln        | Gln        | Val        | Ser       | Asp |

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<210> 344
<211> 516
<212> PRT
<213> Homo sapiens
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|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Met | Ser | Gln | Ser | Thr | Gln | Thr | Asn | Glu | Phe | Leu | Ser | Pro | Glu | Val | Phe |
| 1   |     |     |     | 5   |     |     |     |     | 10  |     |     |     |     | 15  |     |
| Gln | His | Ile | Trp | Asp | Phe | Leu | Glu | Gln | Pro | Ile | Cys | Ser | Val | Gln | Pro |
|     |     |     | 20  |     |     |     |     | 25  |     |     |     |     | 30  |     |     |
| Ile | Asp | Leu | Asn | Phe | Val | Asp | Glu | Pro | Ser | Glu | Asp | Gly | Ala | Thr | Asn |
|     |     | 35  |     |     |     |     | 40  |     |     |     |     | 45  |     |     |     |
| Lys | Ile | Glu | Ile | Ser | Met | Asp | Cys | Ile | Arg | Met | Gln | Asp | Ser | Asp | Leu |
|     | 50  |     |     |     |     | 55  |     |     |     |     | 60  |     |     |     |     |
| Ser | Asp | Pro | Met | Trp | Pro | Gln | Tyr | Thr | Asn | Leu | Gly | Leu | Leu | Asn | Ser |
| 65  |     |     |     |     | 70  |     |     |     |     | 75  |     |     |     |     | 80  |
| Met | Asp | Gln | Gln | Ile | Gln | Asn | Gly | Ser | Ser | Ser | Thr | Ser | Pro | Tyr | Asn |
|     |     |     |     | 85  |     |     |     |     | 90  |     |     |     |     | 95  |     |
| Thr | Asp | His | Ala | Gln | Asn | Ser | Val | Thr | Ala | Pro | Ser | Pro | Tyr | Ala | Gln |
|     |     |     | 100 |     |     |     |     | 105 |     |     |     |     | 110 |     |     |
| Pro | Ser | Ser | Thr | Phe | Asp | Ala | Leu | Ser | Pro | Ser | Pro | Ala | Ile | Pro | Ser |
|     |     |     | 115 |     |     |     | 120 |     |     |     |     | 125 |     |     |     |
| Asn | Thr | Asp | Tyr | Pro | Gly | Pro | His | Ser | Phe | Asp | Val | Ser | Phe | Gln | Gln |
|     | 130 |     |     |     |     | 135 |     |     |     |     | 140 |     |     |     |     |
| Ser | Ser | Thr | Ala | Lys | Ser | Ala | Thr | Trp | Thr | Tyr | Ser | Thr | Glu | Leu | Lys |
| 145 |     |     |     |     | 150 |     |     |     |     | 155 |     |     |     |     | 160 |
| Lys | Leu | Tyr | Cys | Gln | Ile | Ala | Lys | Thr | Cys | Pro | Ile | Gln | Ile | Lys | Val |

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<210> 345
<211> 1800
<212> DNA
<213> Homo sapiens

<400> 345
gcgcctcatt gccactgcag tgactaaagc tgggaagacg ctggtcagtt cacctgcccc 60
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actggttggt ttttaaaca attctgatac aggcgacatc ctcaactgacc gagcaaagat 120
tgacattcgt atcatcactg tgcaccattg gcttctaggc actccagtgg ggtaggagaa 180
ggaggtctga aaccctcgca gagggatctt gccctcattc tttgggtctg aaacactggc 240
agtcgttgga aacaggactc agggataaac cagcgcaatg gattggggga cgctgcacac 300
tttcatcggg ggtgtcaaca aacactccac cagcatcggg aaggtgtgga tcacagtcac 360
ctttattttc cgagtcatga tcctagtggg ggctgcccag gaagtgtggg gtgacgagca 420
agaggacttc gtctgcaaca cactgcaacc gggatgcaaa aatgtgtgct atgaccactt 480
tttcccgggtg tcccacatcc ggctgtgggc cctccagctg atcttcgtct ccaccccagc 540
gctgctgggtg gccatgcatg tggcctacta caggcacgaa accactcgca agttcaggcg 600
aggagagaag aggaatgatt tcaaagacat agaggacatt aaaaagcaca aggttcggat 660
agaggggtcg ctgtgggtgga cgtacaccag cagcatcttt ttccgaatca tctttgaagc 720
agcctttatg tatgtgtttt acttccttta caatgggtac cacctgccct ggggtgtgaa 780
atgtgggatt gaccctgccc ccaaccttgt tgactgcttt atttctaggc caacagagaa 840
gaccgtgttt accattttta tgatttctgc gtctgtgatt tgcattgctgc ttaacgtggc 900
agagtgtgac tacctgctgc tgaaagtgtg ttttaggaga tcaaagagag cacagacgca 960
aaaaaatcac cccaatcatg ccctaaagga gagtaagcag aatgaaatga atgagctgat 1020
ttcagatagt ggtcaaaatg caatcacagg tttcccaagc taaacatttc aaggtaaaat 1080
gtagctgcgt cataaggaga cttctgtctt ctccagaagg caataccaac ctgaaagttc 1140
cttctgtagc ctgaagagtt tgtaaagtac tttcataata aatagacact tgagttaact 1200
ttttgtagga tacttgctcc attcatacac aacgtaatca aatatgtggt ccatctctga 1260
aaacaagaga ctgcttgaca aaggagcatt gcagtcactt tgacaggttc cttttaagtg 1320
gactctctga caaagtgggt actttctgaa aatttatata actgttggtg ataaggaaca 1380
tttatccagg aattgatacg tttattagga aaagatatatt ttataggctt ggatgttttt 1440
agttccgact ttgaatttat ataaagtatt tttataatga ctggtcttcc ttacctggaa 1500
aaacatgcga tgtaggtttt agaattacac cacaagtatc taaatttcca acttacaaag 1560
ggtcctatct tgtaaataatt gttttgcatt gtctgttggc aaatttgtga actgtcatga 1620
tacgcttaag gtgggaaagt gttcattgca caatatatatt ttactgcttt ctgaatgtag 1680
acggaacagt gtggaagcag aaggcttttt taactcatcc gtttggccga tcgttgacaga 1740
ccactgggag atgtggatgt ggttgccctcc ttttgctcgt ccccggtggc taacccttct 1800

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<210> 346

<211> 261

<212> PRT

<213> Homo sapiens

<400> 346

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Met Asp Trp Gly Thr Leu His Thr Phe Ile Gly Gly Val Asn Lys His
 1           5           10           15
Ser Thr Ser Ile Gly Lys Val Trp Ile Thr Val Ile Phe Ile Phe Arg
      20           25           30
Val Met Ile Leu Val Val Ala Ala Gln Glu Val Trp Gly Asp Glu Gln
      35           40           45
Glu Asp Phe Val Cys Asn Thr Leu Gln Pro Gly Cys Lys Asn Val Cys
      50           55           60
Tyr Asp His Phe Phe Pro Val Ser His Ile Arg Leu Trp Ala Leu Gln
      65           70           75           80
Leu Ile Phe Val Ser Thr Pro Ala Leu Leu Val Ala Met His Val Ala
      85           90           95
Tyr Tyr Arg His Glu Thr Thr Arg Lys Phe Arg Arg Gly Glu Lys Arg
      100          105          110
Asn Asp Phe Lys Asp Ile Glu Asp Ile Lys Lys His Lys Val Arg Ile
      115          120          125
Glu Gly Ser Leu Trp Trp Thr Tyr Thr Ser Ser Ile Phe Phe Arg Ile

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|   |     |     |     |     |
|---|-----|-----|-----|-----|
| 130   |     | 135 |     | 140 |
| Ile Phe Glu Ala Ala Phe Met Tyr Val Phe Tyr Phe Leu Tyr Asn Gly |     |     |     |     |
| 145   |     | 150 |     | 155 |
| Tyr His Leu Pro Trp Val Leu Lys Cys Gly Ile Asp Pro Cys Pro Asn |     |     |     |     |
|   | 165 |     | 170 | 175 |
| Leu Val Asp Cys Phe Ile Ser Arg Pro Thr Glu Lys Thr Val Phe Thr |     |     |     |     |
|   | 180 |     | 185 | 190 |
| Ile Phe Met Ile Ser Ala Ser Val Ile Cys Met Leu Leu Asn Val Ala |     |     |     |     |
|   | 195 |     | 200 | 205 |
| Glu Leu Cys Tyr Leu Leu Leu Lys Val Cys Phe Arg Arg Ser Lys Arg |     |     |     |     |
|   | 210 |     | 215 | 220 |
| Ala Gln Thr Gln Lys Asn His Pro Asn His Ala Leu Lys Glu Ser Lys |     |     |     |     |
| 225   |     | 230 |     | 235 |
| Gln Asn Glu Met Asn Glu Leu Ile Ser Asp Ser Gly Gln Asn Ala Ile |     |     |     |     |
|   | 245 |     | 250 | 255 |
| Thr Gly Phe Pro Ser   |     |     |     |     |
| 260   |     |     |     |     |

<210> 347  
 <211> 1740  
 <212> DNA  
 <213> Homo sapiens

<400> 347  
 atgaacaaac tgtatatcgg aaacctcagc gagaacgccg cccctcggga cctagaaagt 60  
 atcttcaagg acgccaagat cccggtgtcg ggacccttcc tggatgaagac tggctacgcg 120  
 ttctgtggact gcccggaaga gagctggggc ctcaaggcca tcgaggcgct ttcaggtaaa 180  
 atagaactgc acgggaaacc catagaagtt gagcactcgg tcccaaaaag gcaaaggatt 240  
 cggaaaacttc agatacgaaa tatcccgccct catttacagt gggagggtgct ggatagttta 300  
 ctagtccagt atggagtggg ggagagctgt gagcaagtga acactgactc ggaaactgca 360  
 gttgtaaatg taacctattc cagtaaggac caagctagac aagcactaga caaactgaat 420  
 ggattttcagt tagagaattt caccttgaaa gttagcctata tccctgatga aacggccgcc 480  
 cagcaaaacc ccttgcagca gccccgaggt cgccggggggc ttgggcagag gggctcctca 540  
 aggcaggggg ctccaggatc cgtatccaag cagaaaccat gtgatttgcc tctgcgcctg 600  
 ctggttccca cccaatttgt tggagccatc ataggaaaag aagggtgccac cattcggaac 660  
 atcaccaaac agaccagtc taaaatcgat gtccaccgta aagaaaatgc gggggctgct 720  
 gagaagtcga ttactatcct ctctactcct gaaggcaact ctgcggcttg taagtctatt 780  
 ctggagatta tgcataagga agctcaagat ataaaattca cagaagagat ccccttgaag 840  
 atttttagctc ataataactt tgttggacgt cttatttggt aagaagggaag aaatcttaaa 900  
 aaaattgagc aagacacaga cactaaaatc acgatatctc cattgcagga attgacgctg 960  
 tataatccag aacgcactat tacagttaaa ggcaatgttg agacatgtgc caaagctgag 1020  
 gaggagatca tgaagaaaat cagggagtct tatgaaaatg atattgcttc tatgaatctt 1080  
 caagcacatt taattcctgg attaaatctg aacgccttgg gtctgttccc acccacttca 1140  
 ggggatgccac ctcccacctc agggccccc tccagccatga ctctcccta cccgcagttt 1200  
 gagcaatcag aaacggagac tgttcatctg tttatcccag ctctatcagt cgggtgccatc 1260  
 atcggaagc agggccagca catcaagcag ctttctcgct ttgctggagc ttcaattaag 1320  
 attgctccag cggaagcacc agatgctaaa gtgaggatgg tgattatcac tggaccacca 1380  
 gaggtcagt tcaaggctca gggaagaatt tatggaaaaa ttaaagaaga aaactttgtt 1440  
 agtcctaaag aagaggtgaa acttgaagct catatcagag tgccatcctt tgctgctggc 1500  
 agagttattg gaaaaggagg caaacgggtg aatgaacttc agaatttgtc aagtgcagaa 1560  
 gttgttgtcc ctctgtacca gacacctgat gagaatgacc aagtggttgt caaaataact 1620  
 ggtcacttct atgcttgcaa gggtgcccag agaaaaattc aggaaattct gactcaggta 1680  
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<210> 348  
 <211> 579  
 <212> PRT  
 <213> Homo sapiens

<400> 348

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Met | Asn | Lys | Leu | Tyr | Ile | Gly | Asn | Leu | Ser | Glu | Asn | Ala | Ala | Pro | Ser |
| 1   |     |     |     | 5   |     |     |     |     | 10  |     |     |     |     | 15  |     |
| Asp | Leu | Glu | Ser | Ile | Phe | Lys | Asp | Ala | Lys | Ile | Pro | Val | Ser | Gly | Pro |
|     |     |     | 20  |     |     |     |     | 25  |     |     |     |     | 30  |     |     |
| Phe | Leu | Val | Lys | Thr | Gly | Tyr | Ala | Phe | Val | Asp | Cys | Pro | Asp | Glu | Ser |
|     |     | 35  |     |     |     |     | 40  |     |     |     |     | 45  |     |     |     |
| Trp | Ala | Leu | Lys | Ala | Ile | Glu | Ala | Leu | Ser | Gly | Lys | Ile | Glu | Leu | His |
|     | 50  |     |     |     |     | 55  |     |     |     |     | 60  |     |     |     |     |
| Gly | Lys | Pro | Ile | Glu | Val | Glu | His | Ser | Val | Pro | Lys | Arg | Gln | Arg | Ile |
| 65  |     |     |     |     | 70  |     |     |     |     | 75  |     |     |     | 80  |     |
| Arg | Lys | Leu | Gln | Ile | Arg | Asn | Ile | Pro | Pro | His | Leu | Gln | Trp | Glu | Val |
|     |     |     |     | 85  |     |     |     | 90  |     |     |     |     |     | 95  |     |
| Leu | Asp | Ser | Leu | Leu | Val | Gln | Tyr | Gly | Val | Val | Glu | Ser | Cys | Glu | Gln |
|     |     |     | 100 |     |     |     |     | 105 |     |     |     |     | 110 |     |     |
| Val | Asn | Thr | Asp | Ser | Glu | Thr | Ala | Val | Val | Asn | Val | Thr | Tyr | Ser | Ser |
|     |     | 115 |     |     |     |     | 120 |     |     |     |     | 125 |     |     |     |
| Lys | Asp | Gln | Ala | Arg | Gln | Ala | Leu | Asp | Lys | Leu | Asn | Gly | Phe | Gln | Leu |
|     | 130 |     |     |     |     | 135 |     |     |     |     | 140 |     |     |     |     |
| Glu | Asn | Phe | Thr | Leu | Lys | Val | Ala | Tyr | Ile | Pro | Asp | Glu | Thr | Ala | Ala |
| 145 |     |     |     |     | 150 |     |     |     |     | 155 |     |     |     |     | 160 |
| Gln | Gln | Asn | Pro | Leu | Gln | Gln | Pro | Arg | Gly | Arg | Arg | Gly | Leu | Gly | Gln |
|     |     |     |     | 165 |     |     |     |     | 170 |     |     |     |     | 175 |     |
| Arg | Gly | Ser | Ser | Arg | Gln | Gly | Ser | Pro | Gly | Ser | Val | Ser | Lys | Gln | Lys |
|     |     |     | 180 |     |     |     |     | 185 |     |     |     |     | 190 |     |     |
| Pro | Cys | Asp | Leu | Pro | Leu | Arg | Leu | Leu | Val | Pro | Thr | Gln | Phe | Val | Gly |
|     |     | 195 |     |     |     |     | 200 |     |     |     |     | 205 |     |     |     |
| Ala | Ile | Ile | Gly | Lys | Glu | Gly | Ala | Thr | Ile | Arg | Asn | Ile | Thr | Lys | Gln |
|     | 210 |     |     |     |     | 215 |     |     |     |     | 220 |     |     |     |     |
| Thr | Gln | Ser | Lys | Ile | Asp | Val | His | Arg | Lys | Glu | Asn | Ala | Gly | Ala | Ala |
| 225 |     |     |     |     | 230 |     |     |     |     | 235 |     |     |     | 240 |     |
| Glu | Lys | Ser | Ile | Thr | Ile | Leu | Ser | Thr | Pro | Glu | Gly | Thr | Ser | Ala | Ala |
|     |     |     | 245 |     |     |     |     |     | 250 |     |     |     |     | 255 |     |
| Cys | Lys | Ser | Ile | Leu | Glu | Ile | Met | His | Lys | Glu | Ala | Gln | Asp | Ile | Lys |
|     |     | 260 |     |     |     |     |     | 265 |     |     |     |     | 270 |     |     |
| Phe | Thr | Glu | Glu | Ile | Pro | Leu | Lys | Ile | Leu | Ala | His | Asn | Asn | Phe | Val |
|     | 275 |     |     |     |     |     | 280 |     |     |     |     | 285 |     |     |     |
| Gly | Arg | Leu | Ile | Gly | Lys | Glu | Gly | Arg | Asn | Leu | Lys | Lys | Ile | Glu | Gln |
|     | 290 |     |     |     |     | 295 |     |     |     |     | 300 |     |     |     |     |
| Asp | Thr | Asp | Thr | Lys | Ile | Thr | Ile | Ser | Pro | Leu | Gln | Glu | Leu | Thr | Leu |
| 305 |     |     |     |     | 310 |     |     |     |     | 315 |     |     |     | 320 |     |
| Tyr | Asn | Pro | Glu | Arg | Thr | Ile | Thr | Val | Lys | Gly | Asn | Val | Glu | Thr | Cys |
|     |     |     |     | 325 |     |     |     |     | 330 |     |     |     |     | 335 |     |
| Ala | Lys | Ala | Glu | Glu | Glu | Ile | Met | Lys | Lys | Ile | Arg | Glu | Ser | Tyr | Glu |
|     |     | 340 |     |     |     |     |     | 345 |     |     |     |     | 350 |     |     |
| Asn | Asp | Ile | Ala | Ser | Met | Asn | Leu | Gln | Ala | His | Leu | Ile | Pro | Gly | Leu |
|     |     | 355 |     |     |     |     | 360 |     |     |     |     |     | 365 |     |     |

Asn Leu Asn Ala Leu Gly Leu Phe Pro Pro Thr Ser Gly Met Pro Pro  
 370 375 380  
 Pro Thr Ser Gly Pro Pro Ser Ala Met Thr Pro Pro Tyr Pro Gln Phe  
 385 390 395 400  
 Glu Gln Ser Glu Thr Glu Thr Val His Leu Phe Ile Pro Ala Leu Ser  
 405 410 415  
 Val Gly Ala Ile Ile Gly Lys Gln Gly Gln His Ile Lys Gln Leu Ser  
 420 425 430  
 Arg Phe Ala Gly Ala Ser Ile Lys Ile Ala Pro Ala Glu Ala Pro Asp  
 435 440 445  
 Ala Lys Val Arg Met Val Ile Ile Thr Gly Pro Pro Glu Ala Gln Phe  
 450 455 460  
 Lys Ala Gln Gly Arg Ile Tyr Gly Lys Ile Lys Glu Glu Asn Phe Val  
 465 470 475 480  
 Ser Pro Lys Glu Glu Val Lys Leu Glu Ala His Ile Arg Val Pro Ser  
 485 490 495  
 Phe Ala Ala Gly Arg Val Ile Gly Lys Gly Gly Lys Thr Val Asn Glu  
 500 505 510  
 Leu Gln Asn Leu Ser Ser Ala Glu Val Val Val Pro Arg Asp Gln Thr  
 515 520 525  
 Pro Asp Glu Asn Asp Gln Val Val Val Lys Ile Thr Gly His Phe Tyr  
 530 535 540  
 Ala Cys Gln Val Ala Gln Arg Lys Ile Gln Glu Ile Leu Thr Gln Val  
 545 550 555 560  
 Lys Gln His Gln Gln Gln Lys Ala Leu Gln Ser Gly Pro Pro Gln Ser  
 565 570 575  
 Arg Arg Lys

<210> 349  
 <211> 207  
 <212> DNA  
 <213> Homo sapiens

<400> 349  
 atgtggcagc ccctcttctt caagtggctc ttgtcctggt gccctgggag ttctcaaatt 60  
 gctgcagcag cctccaccca gcctgaggat gacatcaata cacagaggaa gaagagtcag 120  
 gaaaagatga gagaagttac agactctcct gggcgacccc gagagcttac cattcctcag 180  
 acttcttcac atggtgctaa cagattt 207

<210> 350  
 <211> 69  
 <212> PRT  
 <213> Homo sapiens

<400> 350  
 Met Trp Gln Pro Leu Phe Phe Lys Trp Leu Leu Ser Cys Cys Pro Gly  
 1 5 10 15  
 Ser Ser Gln Ile Ala Ala Ala Ala Ser Thr Gln Pro Glu Asp Asp Ile  
 20 25 30  
 Asn Thr Gln Arg Lys Lys Ser Gln Glu Lys Met Arg Glu Val Thr Asp  
 35 40 45  
 Ser Pro Gly Arg Pro Arg Glu Leu Thr Ile Pro Gln Thr Ser Ser His

50  
Gly Ala Asn Arg Phe  
65

55

60

<210> 351  
<211> 1012  
<212> DNA  
<213> Homo sapiens

<400> 351  
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catcacacgg ccgcgtccga taacttccag ctgtcccagg gtgggcaggg attcgccatt 120  
ccgatcgggc aggcgatggc gatcgcgggc cagatcaagc tccccaccgt tcatatcggg 180  
cctaccgcct tctcggctt ggggtgtgtc gacaacaacg gcaacggcgc acgagtccaa 240  
cgcggtggtc ggagcgctcc ggcggcaagt ctcggcacat ccaccggcga cgtgatcacc 300  
gcggtcgacg gcgctccgat caactcggcc accgcgatgg cggacgcgct taacgggcat 360  
catcccggtg acgtcatctc ggtgacctgg caaaccaagt cgggcggcac gcgtacaggg 420  
aacgtgacat tggccgaggg acccccggcc gaattcatgg attgggggac gctgcacact 480  
ttcatcgggg gtgtcaacaa acactccacc agcatcggga aggtgtggat cacagtcatc 540  
tttattttcc gagtcatgat cctcgtgggt gctgcccagg aagtgtgggg tgacgagcaa 600  
gaggacttcg tctgcaaacac actgcaaccg ggatgcaaaa atgtgtgcta tgaccacttt 660  
tccccgggtg cccacatccg gctgtggggc ctccagctga tcttcgtctc cccccagcg 720  
ctgctgggtg ccatgcatgt ggctactac aggcacgaaa ccactcgcaa gttcaggcga 780  
ggagagaaga ggaatgattt caaagacata gaggacatta aaaagcagaa ggttcggata 840  
gaggggtgac tcgagcacca ccaccaccac cactgagatc cggtctgtaa caaagcccga 900  
aaggaagctg agttggctgc tgccaccgct gagcaataac tagcataacc ccttggggcc 960  
tctaaacggg tcttgagggg ttttttgctg aaaggaggaa ctatatccgg at 1012

<210> 352  
<211> 267  
<212> PRT  
<213> Homo sapiens

<400> 352  
Met His His His His His His Thr Ala Ala Ser Asp Asn Phe Gln Leu  
1 5 10 15  
Ser Gln Gly Gly Gln Gly Phe Ala Ile Pro Ile Gly Gln Ala Met Ala  
20 25 30  
Ile Ala Gly Gln Ile Lys Leu Pro Thr Val His Ile Gly Pro Thr Ala  
35 40 45  
Phe Leu Gly Leu Gly Val Val Asp Asn Asn Gly Asn Gly Ala Arg Val  
50 55 60  
Gln Arg Val Val Gly Ser Ala Pro Ala Ala Ser Leu Gly Ile Ser Thr  
65 70 75 80  
Gly Asp Val Ile Thr Ala Val Asp Gly Ala Pro Ile Asn Ser Ala Thr  
85 90 95  
Ala Met Ala Asp Ala Leu Asn Gly His His Pro Gly Asp Val Ile Ser  
100 105 110  
Val Thr Trp Gln Thr Lys Ser Gly Gly Thr Arg Thr Gly Asn Val Thr  
115 120 125  
Leu Ala Glu Gly Pro Pro Ala Glu Phe Met Asp Trp Gly Thr Leu His  
130 135 140  
Thr Phe Ile Gly Gly Val Asn Lys His Ser Thr Ser Ile Gly Lys Val

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145          150          155          160
Trp Ile Thr Val Ile Phe Ile Phe Arg Val Met Ile Leu Val Val Ala
          165          170          175
Ala Gln Glu Val Trp Gly Asp Glu Gln Glu Asp Phe Val Cys Asn Thr
          180          185          190
Leu Gln Pro Gly Cys Lys Asn Val Cys Tyr Asp His Phe Phe Pro Val
          195          200          205
Ser His Ile Arg Leu Trp Ala Leu Gln Leu Ile Phe Val Ser Thr Pro
          210          215          220
Ala Leu Leu Val Ala Met His Val Ala Tyr Tyr Arg His Glu Thr Thr
225          230          235          240
Arg Lys Phe Arg Arg Gly Glu Lys Arg Asn Asp Phe Lys Asp Ile Glu
          245          250          255
Asp Ile Lys Lys Gln Lys Val Arg Ile Glu Gly
          260          265

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<210> 353
<211> 900
<212> DNA
<213> Homo sapiens

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<400> 353
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accgttcata tcgggcctac cgcccttcctc ggcttggttg ttgtcgacaa caacggcaac 180
ggcgcacgag tccaacgcgt ggtcgggagc gctccggcgg caagtctcgg catctccacc 240
ggcgacgtga tcaccgcggt cgacggcgct ccgatcaact cggccaccgc gatggcggac 300
gcgcttaacg ggcattcatc cgggtgacgt atctcgggtga cctggcaaac caagtcgggc 360
ggcacgcgta cagggaaact gacattggcc gagggacccc cggccgaatt ccacgaaacc 420
actcgcaagt tcaggcgagg agagaagagg aatgatttca aagacataga ggacattaaa 480
aagcagaagg ttcggtataga ggggtcgctg tgggtggacgt acaccagcag catctttttc 540
cgaatcatct ttgaagcagc ctttatgtat gtgttttact tcctttacaa tgggtaccac 600
ctgccctggg tgttgaaatg tgggattgac ccctgcccc aacctgttga ctgctttatt 660
tctaggccaa cagagaagac cgtgtttacc atttttatga tttctgcgtc tgtgatttgc 720
atgctgctta acgtggcaga gttgtgctac ctgctgctga aagtgtgttt taggagatca 780
aagagagcac agacgcaaaa aaatcacccc aatcatgccc taaaggagag taagcagaat 840
gaaatgaatg agctgatttc agatagtggg caaaatgcaa tcacagggtt cccaagctaa 900

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<210> 354
<211> 299
<212> PRT
<213> Homo sapiens

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<400> 354
Met His His His His His His Thr Ala Ala Ser Asp Asn Phe Gln Leu
  1          5          10          15
Ser Gln Gly Gly Gln Gly Phe Ala Ile Pro Ile Gly Gln Ala Met Ala
          20          25          30
Ile Ala Gly Gln Ile Lys Leu Pro Thr Val His Ile Gly Pro Thr Ala
          35          40          45
Phe Leu Gly Leu Gly Val Val Asp Asn Asn Gly Asn Gly Ala Arg Val
          50          55          60

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Gln Arg Val Val Gly Ser Ala Pro Ala Ala Ser Leu Gly Ile Ser Thr  
 65 70 75 80  
 Gly Asp Val Ile Thr Ala Val Asp Gly Ala Pro Ile Asn Ser Ala Thr  
 85 90 95  
 Ala Met Ala Asp Ala Leu Asn Gly His His Pro Gly Asp Val Ile Ser  
 100 105 110  
 Val Thr Trp Gln Thr Lys Ser Gly Gly Thr Arg Thr Gly Asn Val Thr  
 115 120 125  
 Leu Ala Glu Gly Pro Pro Ala Glu Phe His Glu Thr Thr Arg Lys Phe  
 130 135 140  
 Arg Arg Gly Glu Lys Arg Asn Asp Phe Lys Asp Ile Glu Asp Ile Lys  
 145 150 155 160  
 Lys Gln Lys Val Arg Ile Glu Gly Ser Leu Trp Trp Thr Tyr Thr Ser  
 165 170 175  
 Ser Ile Phe Phe Arg Ile Ile Phe Glu Ala Ala Phe Met Tyr Val Phe  
 180 185 190  
 Tyr Phe Leu Tyr Asn Gly Tyr His Leu Pro Trp Val Leu Lys Cys Gly  
 195 200 205  
 Ile Asp Pro Cys Pro Asn Leu Val Asp Cys Phe Ile Ser Arg Pro Thr  
 210 215 220  
 Glu Lys Thr Val Phe Thr Ile Phe Met Ile Ser Ala Ser Val Ile Cys  
 225 230 235 240  
 Met Leu Leu Asn Val Ala Glu Leu Cys Tyr Leu Leu Leu Lys Val Cys  
 245 250 255  
 Phe Arg Arg Ser Lys Arg Ala Gln Thr Gln Lys Asn His Pro Asn His  
 260 265 270  
 Ala Leu Lys Glu Ser Lys Gln Asn Glu Met Asn Glu Leu Ile Ser Asp  
 275 280 285  
 Ser Gly Gln Asn Ala Ile Thr Gly Phe Pro Ser  
 290 295

<210> 355

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 355

ggagtacagc ttcaagacaa tggg

24

<210> 356

<211> 31

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 356

ccatgggaat tcattataat aattttgttc c

31

<210> 357  
 <211> 920  
 <212> PRT  
 <213> Homo sapiens

<400> 357

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Met | Gln | His | His | His | His | His | His | Gly | Val | Gln | Leu | Gln | Asp | Asn | Gly |
| 1   |     |     |     | 5   |     |     |     |     | 10  |     |     |     |     | 15  |     |
| Tyr | Asn | Gly | Leu | Leu | Ile | Ala | Ile | Asn | Pro | Gln | Val | Pro | Glu | Asn | Gln |
|     |     |     | 20  |     |     |     |     | 25  |     |     |     |     | 30  |     |     |
| Asn | Leu | Ile | Ser | Asn | Ile | Lys | Glu | Met | Ile | Thr | Glu | Ala | Ser | Phe | Tyr |
|     |     | 35  |     |     |     |     | 40  |     |     |     |     | 45  |     |     |     |
| Leu | Phe | Asn | Ala | Thr | Lys | Arg | Val | Phe | Phe | Arg | Asn | Ile | Lys | Ile |     |
|     | 50  |     |     |     |     | 55  |     |     |     | 60  |     |     |     |     |     |
| Leu | Ile | Pro | Ala | Thr | Trp | Lys | Ala | Asn | Asn | Asn | Ser | Lys | Ile | Lys | Gln |
| 65  |     |     |     |     | 70  |     |     |     |     | 75  |     |     |     |     | 80  |
| Glu | Ser | Tyr | Glu | Lys | Ala | Asn | Val | Ile | Val | Thr | Asp | Trp | Tyr | Gly | Ala |
|     |     |     |     | 85  |     |     |     |     | 90  |     |     |     |     | 95  |     |
| His | Gly | Asp | Asp | Pro | Tyr | Thr | Leu | Gln | Tyr | Arg | Gly | Cys | Gly | Lys | Glu |
|     |     |     | 100 |     |     |     |     | 105 |     |     |     |     | 110 |     |     |
| Gly | Lys | Tyr | Ile | His | Phe | Thr | Pro | Asn | Phe | Leu | Leu | Asn | Asp | Asn | Leu |
|     |     | 115 |     |     |     |     | 120 |     |     |     |     | 125 |     |     |     |
| Thr | Ala | Gly | Tyr | Gly | Ser | Arg | Gly | Arg | Val | Phe | Val | His | Glu | Trp | Ala |
|     | 130 |     |     |     |     | 135 |     |     |     |     | 140 |     |     |     |     |
| His | Leu | Arg | Trp | Gly | Val | Phe | Asp | Glu | Tyr | Asn | Asn | Asp | Lys | Pro | Phe |
| 145 |     |     |     |     | 150 |     |     |     |     | 155 |     |     |     |     | 160 |
| Tyr | Ile | Asn | Gly | Gln | Asn | Gln | Ile | Lys | Val | Thr | Arg | Cys | Ser | Ser | Asp |
|     |     |     |     | 165 |     |     |     |     | 170 |     |     |     |     | 175 |     |
| Ile | Thr | Gly | Ile | Phe | Val | Cys | Glu | Lys | Gly | Pro | Cys | Pro | Gln | Glu | Asn |
|     |     |     | 180 |     |     |     |     | 185 |     |     |     |     | 190 |     |     |
| Cys | Ile | Ile | Ser | Lys | Leu | Phe | Lys | Glu | Gly | Cys | Thr | Phe | Ile | Tyr | Asn |
|     |     | 195 |     |     |     |     | 200 |     |     |     |     | 205 |     |     |     |
| Ser | Thr | Gln | Asn | Ala | Thr | Ala | Ser | Ile | Met | Phe | Met | Gln | Ser | Leu | Ser |
|     |     | 210 |     |     |     | 215 |     |     |     |     | 220 |     |     |     |     |
| Ser | Val | Val | Glu | Phe | Cys | Asn | Ala | Ser | Thr | His | Asn | Gln | Glu | Ala | Pro |
| 225 |     |     |     |     | 230 |     |     |     |     | 235 |     |     |     |     | 240 |
| Asn | Leu | Gln | Asn | Gln | Met | Cys | Ser | Leu | Arg | Ser | Ala | Trp | Asp | Val | Ile |
|     |     |     |     | 245 |     |     |     |     | 250 |     |     |     |     | 255 |     |
| Thr | Asp | Ser | Ala | Asp | Phe | His | His | Ser | Phe | Pro | Met | Asn | Gly | Thr | Glu |
|     |     |     | 260 |     |     |     |     | 265 |     |     |     |     | 270 |     |     |
| Leu | Pro | Pro | Pro | Pro | Thr | Phe | Ser | Leu | Val | Glu | Ala | Gly | Asp | Lys | Val |
|     |     | 275 |     |     |     |     | 280 |     |     |     |     | 285 |     |     |     |
| Val | Cys | Leu | Val | Leu | Asp | Val | Ser | Ser | Lys | Met | Ala | Glu | Ala | Asp | Arg |
|     |     | 290 |     |     |     | 295 |     |     |     |     | 300 |     |     |     |     |
| Leu | Leu | Gln | Leu | Gln | Gln | Ala | Ala | Glu | Phe | Tyr | Leu | Met | Gln | Ile | Val |
| 305 |     |     |     |     | 310 |     |     |     |     | 315 |     |     |     |     | 320 |
| Glu | Ile | His | Thr | Phe | Val | Gly | Ile | Ala | Ser | Phe | Asp | Ser | Lys | Gly | Glu |
|     |     |     |     | 325 |     |     |     | 330 |     |     |     |     |     | 335 |     |
| Ile | Arg | Ala | Gln | Leu | His | Gln | Ile | Asn | Ser | Asn | Asp | Asp | Arg | Lys | Leu |
|     |     |     | 340 |     |     |     | 345 |     |     |     |     |     | 350 |     |     |
| Leu | Val | Ser | Tyr | Leu | Pro | Thr | Thr | Val | Ser | Ala | Lys | Thr | Asp | Ile | Ser |
|     |     | 355 |     |     |     |     | 360 |     |     |     |     | 365 |     |     |     |
| Ile | Cys | Ser | Gly | Leu | Lys | Lys | Gly | Phe | Glu | Val | Val | Glu | Lys | Leu | Asn |
|     |     | 370 |     |     |     | 375 |     |     |     |     |     | 380 |     |     |     |

|         |     |     |     |         |         |     |     |     |         |         |     |     |     |         |
|---------|-----|-----|-----|---------|---------|-----|-----|-----|---------|---------|-----|-----|-----|---------|
| Gly 385 | Lys | Ala | Tyr | Gly     | Ser 390 | Val | Met | Ile | Leu     | Val 395 | Thr | Ser | Gly | Asp 400 |
| Lys     | Leu | Leu | Gly | Asn 405 | Cys     | Leu | Pro | Thr | Val 410 | Leu     | Ser | Ser | Gly | Ser 415 |
| Ile     | His | Ser | Ile | Ala 420 | Leu     | Gly | Ser | Ser | Ala 425 | Ala     | Pro | Asn | Leu | Glu 430 |
| Leu     | Ser | Arg | Leu | Thr 435 | Gly     | Gly | Leu | Lys | Phe 440 | Phe     | Val | Pro | Asp | Ile 445 |
| Asn     | Ser | Asn | Ser | Met 450 | Ile     | Asp | Ala | Phe | Ser 455 | Arg     | Ile | Ser | Ser | Gly 460 |
| Gly 465 | Asp | Ile | Phe | Gln 470 | Gln     | His | Ile | Gln | Leu 475 | Glu     | Ser | Thr | Gly | Glu 480 |
| Val     | Lys | Pro | His | His 485 | Gln     | Leu | Lys | Asn | Thr 490 | Val     | Thr | Val | Asp | Asn 495 |
| Val     | Gly | Asn | Asp | Thr 500 | Met     | Phe | Leu | Val | Thr 505 | Trp     | Gln | Ala | Ser | Gly 510 |
| Pro     | Glu | Ile | Ile | Leu 515 | Phe     | Asp | Pro | Asp | Gly 520 | Arg     | Lys | Tyr | Tyr | Thr 525 |
| Asn     | Phe | Ile | Thr | Asn 530 | Leu     | Thr | Phe | Arg | Thr 535 | Ala     | Ser | Leu | Trp | Ile 540 |
| Gly 545 | Thr | Ala | Lys | Pro 550 | Gly     | His | Trp | Thr | Tyr 555 | Thr     | Leu | Asn | Asn | Thr 560 |
| His     | Ser | Leu | Gln | Ala 565 | Leu     | Lys | Val | Thr | Val 570 | Thr     | Ser | Arg | Ala | Ser 575 |
| Ser     | Ala | Val | Pro | Ala 580 | Thr     | Val | Glu | Ala | Phe 585 | Val     | Glu | Arg | Asp | Ser 590 |
| Leu     | His | Phe | Pro | His 595 | Pro     | Val | Met | Ile | Tyr 600 | Ala     | Asn | Val | Lys | Gln 605 |
| Phe     | Tyr | Pro | Ile | Leu 610 | Asn     | Ala | Thr | Val | Thr 615 | Ala     | Thr | Val | Glu | Pro 620 |
| Thr 625 | Gly | Asp | Pro | Val 630 | Thr     | Leu | Arg | Leu | Leu 635 | Asp     | Asp | Gly | Ala | Gly 640 |
| Asp     | Val | Ile | Lys | Asn 645 | Asp     | Gly | Ile | Tyr | Ser 650 | Arg     | Tyr | Phe | Phe | Ser 655 |
| Ala     | Ala | Asn | Gly | Arg 660 | Tyr     | Ser | Leu | Lys | Val 665 | His     | Val | Asn | His | Ser 670 |
| Ser     | Ile | Ser | Thr | Pro 675 | Ala     | His | Ser | Ile | Pro 680 | Gly     | Ser | His | Ala | Met 685 |
| Val     | Pro | Gly | Tyr | Thr 690 | Ala     | Asn | Gly | Asn | Ile 695 | Gln     | Met | Asn | Ala | Pro 700 |
| Lys 705 | Ser | Val | Gly | Arg 710 | Asn     | Glu | Glu | Glu | Arg 715 | Lys     | Trp | Gly | Phe | Ser 720 |
| Val     | Ser | Ser | Gly | Gly 725 | Ser     | Phe | Ser | Val | Leu 730 | Gly     | Val | Pro | Ala | Gly 735 |
| His     | Pro | Asp | Val | Phe 740 | Pro     | Pro | Cys | Lys | Ile 745 | Ile     | Asp | Leu | Glu | Ala 750 |
| Lys     | Val | Glu | Glu | Glu 755 | Leu     | Thr | Leu | Ser | Trp 760 | Thr     | Ala | Pro | Gly | Glu 765 |
| Phe     | Asp | Gln | Gly | Gln 770 | Ala     | Thr | Ser | Tyr | Glu 775 | Ile     | Arg | Met | Ser | Lys 780 |
| Leu 785 | Gln | Asn | Ile | Gln 790 | Asp     | Phe | Asn | Asn | Ala 795 | Ile     | Leu | Val | Asn | Thr 800 |
| Ser     | Lys | Arg | Asn | Pro 805 | Gln     | Gln | Ala | Gly | Ile 810 | Arg     | Glu | Ile | Phe | Thr 815 |



Ser Pro Gln Ile Ser Thr Asn Gly Pro Glu His Gln Pro Asn Gly Glu  
                     820                    825                    830  
 Thr His Glu Ser His Arg Ile Tyr Val Ala Ile Arg Ala Met Asp Arg  
                     835                    840                    845  
 Asn Ser Leu Gln Ser Ala Val Ser Asn Ile Ala Gln Ala Pro Leu Phe  
                     850                    855                    860  
 Ile Pro Pro Asn Ser Asp Pro Val Pro Ala Arg Asp Tyr Leu Ile Leu  
 865                    870                    875                    880  
 Lys Gly Val Leu Thr Ala Met Gly Leu Ile Gly Ile Ile Cys Leu Ile  
                     885                    890                    895  
 Ile Val Val Thr His His Thr Leu Ser Arg Lys Lys Arg Ala Asp Lys  
                     900                    905                    910  
 Lys Glu Asn Gly Thr Lys Leu Leu  
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<210> 358

<211> 2773

<212> DNA

<213> Homo sapiens

<400> 358

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 agaaatataa agatttttaac acctgccaca tggaaagcta ataataacag caaaataaaa 240  
 caagaatcat atgaaaaggc aaatgtcata gtgactgact ggtatggggc acatggagat 300  
 gatccataca ccttacaata cagagggtgt ggaaaagagg gaaaatacat tcatttcaca 360  
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 acagttgagc cagagactgg agatcctgtt acgctgagac tccttgatga tggagcaggt 1920

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gctgatgtta taaaaaatga tggaattttac tgcaggtatt ttttctcctt tgctgcaaatt 1980
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tctattccag ggagtcacgc tatgtatgta ccagggtaca cagcaaacgg taatattcag 2100
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<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 359

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<210> 360

<211> 33

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 360

cgccagaatt catcaaaca atctgttagc acc 33

<210> 361

<211> 77

<212> PRT

<213> Homo sapiens

<400> 361

Met Gln His His His His His His Trp Gln Pro Leu Phe Phe Lys Trp

1 5 10 15

Leu Leu Ser Cys Cys Pro Gly Ser Ser Gln Ile Ala Ala Ala Ala Ser

20 25 30

Thr Gln Pro Glu Asp Asp Ile Asn Thr Gln Arg Lys Lys Ser Gln Glu

35 40 45

Lys Met Arg Glu Val Thr Asp Ser Pro Gly Arg Pro Arg Glu Leu Thr

50 55 60

Ile Pro Gln Thr Ser Ser His Gly Ala Asn Arg Phe Val

65 70 75

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 <211> 244  
 <212> DNA  
 <213> Homo sapiens

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 aatacacaga ggaagaagag tcaggaaaag atgagagaag ttacagactc tcctgggcga 180  
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 attc 244

<210> 363  
 <211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 363  
 Met Trp Gln Pro Leu Phe Phe Lys Trp Leu Leu Ser Cys Cys Pro Gly  
 1 5 10 15  
 Ser Ser Gln Ile  
 20

<210> 364  
 <211> 60  
 <212> DNA  
 <213> Homo sapiens

<400> 364  
 atgtggcagc cctctttctt caagtggctc ttgtcctggt gccctggggag ttctcaaatt 60

<210> 365  
 <211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 365  
 Gly Ser Ser Gln Ile Ala Ala Ala Ala Ser Thr Gln Pro Glu Asp Asp  
 1 5 10 15  
 Ile Asn Thr Gln  
 20

<210> 366  
 <211> 60  
 <212> DNA  
 <213> Homo sapiens

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<210> 367  
 <211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 367  
 Lys Pro Gly His Trp Thr Tyr Thr Leu Asn Asn Thr His His Ser Leu  
 1 5 10 15  
 Gln Ala Leu Lys  
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<210> 368  
 <211> 2343  
 <212> DNA  
 <213> Homo sapiens

<400> 368  
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 gcgccgcgcc tctgaggcgc agcatgtgaa gcggagacgg catccagtgg ggggcgagcc 180  
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 atggatgaaa ggattcagag gaaagtagag aaactagagc aacaatgtca gaaagaagcc 420  
 aaggaatttg ccaagaaggt acaagagctg cagaaaagca atcaggttgc cttccaacat 480  
 ttccaagaac tagatgagca ctttagctat gtagcaacta aagtctgtca cttgggagac 540  
 cagtttagagg gggtaaacc acccagacaa cgggcagtgg aggctcagaa attgatgaaa 600  
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 gggagaagaat tggatacagt tttgatggaa cttggagtag gttttcatcg acttatctat 2040

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tcaggagaac aacttgctaa tctggacaag aatatacttc actccttcgt acaacttcgt 2280
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att 2343

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<210> 369
<211> 708
<212> PRT
<213> Homo sapiens

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Tyr Ile Glu Arg Leu Val Trp Arg Thr Pro Gly Gly Gly Ser Arg Gly
20     25     30
Gly Pro Glu Ala Phe Asp Pro Lys Arg Leu Leu Glu Glu Phe Val Asn
35     40     45
His Ile Gln Glu Leu Gln Ile Met Asp Glu Arg Ile Gln Arg Lys Val
50     55     60
Glu Lys Leu Glu Gln Gln Cys Gln Lys Glu Ala Lys Glu Phe Ala Lys
65     70     75     80
Lys Val Gln Glu Leu Gln Lys Ser Asn Gln Val Ala Phe Gln His Phe
85     90     95
Gln Glu Leu Asp Glu His Ile Ser Tyr Val Ala Thr Lys Val Cys His
100    105    110
Leu Gly Asp Gln Leu Glu Gly Val Asn Thr Pro Arg Gln Arg Ala Val
115    120    125
Glu Ala Gln Lys Leu Met Lys Tyr Phe Asn Glu Phe Leu Asp Gly Glu
130    135    140
Leu Lys Ser Asp Val Phe Thr Asn Ser Glu Lys Ile Lys Glu Ala Ala
145    150    155    160
Asp Ile Ile Gln Lys Leu His Leu Ile Ala Gln Glu Leu Pro Phe Asp
165    170    175
Arg Phe Ser Glu Val Lys Ser Lys Ile Ala Ser Lys Tyr His Asp Leu
180    185    190
Glu Cys Gln Leu Ile Gln Glu Phe Thr Ser Ala Gln Arg Arg Gly Glu
195    200    205
Ile Ser Arg Met Arg Glu Val Ala Ala Val Leu Leu His Phe Lys Gly
210    215    220
Tyr Ser His Cys Val Asp Val Tyr Ile Lys Gln Cys Gln Glu Gly Ala
225    230    235    240
Tyr Leu Arg Asn Asp Ile Phe Glu Asp Ala Gly Ile Leu Cys Gln Arg
245    250    255
Val Asn Lys Gln Val Gly Asp Ile Phe Ser Asn Pro Glu Thr Val Leu
260    265    270
Ala Lys Leu Ile Gln Asn Val Phe Glu Ile Lys Leu Gln Ser Phe Val
275    280    285
Lys Glu Gln Leu Glu Glu Cys Arg Lys Ser Asp Ala Glu Gln Tyr Leu
290    295    300
Lys Asn Leu Tyr Asp Leu Tyr Thr Arg Thr Thr Asn Leu Ser Ser Lys
305    310    315    320
Leu Met Glu Phe Asn Leu Gly Thr Asp Lys Gln Thr Phe Leu Ser Lys

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Leu Ile Lys Ser Ile Phe Ile Ser Tyr Leu Glu Asn Tyr Ile Glu Val  
 325 340 345 350  
 Glu Thr Gly Tyr Leu Lys Ser Arg Ser Ala Met Ile Leu Gln Arg Tyr  
 355 360 365  
 Tyr Asp Ser Lys Asn His Gln Lys Arg Ser Ile Gly Thr Gly Gly Ile  
 370 375 380  
 Gln Asp Leu Lys Glu Arg Ile Arg Gln Arg Thr Asn Leu Pro Leu Gly  
 385 390 395 400  
 Pro Ser Ile Asp Thr His Gly Glu Thr Phe Leu Ser Gln Glu Val Val  
 405 410 415  
 Val Asn Leu Leu Gln Glu Thr Lys Gln Ala Phe Glu Arg Cys His Arg  
 420 425 430  
 Leu Ser Asp Pro Ser Asp Leu Pro Arg Asn Ala Phe Arg Ile Phe Thr  
 435 440 445  
 Ile Leu Val Glu Phe Leu Cys Ile Glu His Ile Asp Tyr Ala Leu Glu  
 450 455 460  
 Thr Gly Leu Ala Gly Ile Pro Ser Ser Asp Ser Arg Asn Ala Asn Leu  
 465 470 475 480  
 Tyr Phe Leu Asp Val Val Gln Gln Ala Asn Thr Ile Phe His Leu Phe  
 485 490 495  
 Asp Lys Gln Phe Asn Asp His Leu Met Pro Leu Ile Ser Ser Ser Pro  
 500 505 510  
 Lys Leu Ser Glu Cys Leu Gln Lys Lys Lys Glu Ile Ile Glu Gln Met  
 515 520 525  
 Glu Met Lys Leu Asp Thr Gly Ile Asp Arg Thr Leu Asn Cys Met Ile  
 530 535 540  
 Gly Gln Met Lys His Ile Leu Ala Ala Glu Gln Lys Lys Thr Asp Phe  
 545 550 555 560  
 Lys Pro Glu Asp Glu Asn Asn Val Leu Ile Gln Tyr Thr Asn Ala Cys  
 565 570 575  
 Val Lys Val Cys Ala Tyr Val Arg Lys Gln Val Glu Lys Ile Lys Asn  
 580 585 590  
 Ser Met Asp Gly Lys Asn Val Asp Thr Val Leu Met Glu Leu Gly Val  
 595 600 605  
 Arg Phe His Arg Leu Ile Tyr Glu His Leu Gln Gln Tyr Ser Tyr Ser  
 610 615 620  
 Cys Met Gly Gly Met Leu Ala Ile Cys Asp Val Ala Glu Tyr Arg Lys  
 625 630 635 640  
 Cys Ala Lys Asp Phe Lys Ile Pro Met Val Leu His Leu Phe Asp Thr  
 645 650 655  
 Leu His Ala Leu Cys Asn Leu Leu Val Val Ala Pro Asp Asn Leu Lys  
 660 665 670  
 Gln Val Cys Ser Gly Glu Gln Leu Ala Asn Leu Asp Lys Asn Ile Leu  
 675 680 685  
 His Ser Phe Val Gln Leu Arg Ala Asp Tyr Arg Ser Ala Arg Leu Ala  
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 Arg His Phe Ser  
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&lt;210&gt; 370

&lt;211&gt; 60

&lt;212&gt; DNA

<213> Homo sapiens

<400> 370

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<210> 371

<211> 60

<212> DNA

<213> Homo sapiens

<400> 371

agtagaattt cctctggaac tggagacatt ttccagcaac atattcagct tgaaagtaca 60

<210> 372

<211> 60

<212> DNA

<213> Homo sapiens

<400> 372

ccagagactg gagatcctgt tacgctgaga ctccttgatg atggagcagg tgctgatgtt 60

<210> 373

<211> 60

<212> DNA

<213> Homo sapiens

<400> 373

ttacagtctg ctgtatctaa cattgcccag gcgcctctgt ttattccccc caattctgat 60

<210> 374

<211> 60

<212> DNA

<213> Homo sapiens

<400> 374

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<210> 375

<211> 60

<212> DNA

<213> Homo sapiens

<400> 375

aaaaacacag tgactgtgga taatactgtg ggcaacgaca ctatgtttct agttacgtgg 60

<210> 376

<211> 20

<212> PRT

<213> Homo sapiens

<400> 376

Leu Gln Ser Ala Val Ser Asn Ile Ala Gln Ala Pro Leu Phe Ile Pro  
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 Pro Asn Ser Asp  
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<210> 377

<211> 20

<212> PRT

<213> Homo sapiens

<400> 377

Val Asn His Ser Pro Ser Ile Ser Thr Pro Ala His Ser Ile Pro Gly  
 1 5 10 15  
 Ser His Ala Met  
 20

<210> 378

<211> 20

<212> PRT

<213> Homo sapiens

<400> 378

Pro Glu Thr Gly Asp Pro Val Thr Leu Arg Leu Leu Asp Asp Gly Ala  
 1 5 10 15  
 Gly Ala Asp Val  
 20

<210> 379

<211> 20

<212> PRT

<213> Homo sapiens

<400> 379

Ala Val Pro Pro Ala Thr Val Glu Ala Phe Val Glu Arg Asp Ser Leu  
 1 5 10 15  
 His Phe Pro His  
 20

<210> 380

<211> 20

<212> PRT

<213> Homo sapiens

<400> 380

Ser Arg Ile Ser Ser Gly Thr Gly Asp Ile Phe Gln Gln His Ile Gln  
 1 5 10 15  
 Leu Glu Ser Thr



20

&lt;210&gt; 381

&lt;211&gt; 20

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 381

Lys Asn Thr Val Thr Val Asp Asn Thr Val Gly Asn Asp Thr Met Phe  
 1 5 10 15

Leu Val Thr Trp  
 20

&lt;210&gt; 382

&lt;211&gt; 20

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 382

Lys Pro Gly His Trp Thr Tyr Thr Leu Asn Asn Thr His His Ser Leu  
 1 5 10 15

Gln Ala Leu Lys  
 20

&lt;210&gt; 383

&lt;211&gt; 29

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; PCR primer

&lt;400&gt; 383

cggcgaattc atggattggg ggacgctgc

29

&lt;210&gt; 384

&lt;211&gt; 35

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; PCR primer

&lt;400&gt; 384

cggcctcgag tcacccctct atccgaacct tctgc

35

&lt;210&gt; 385

&lt;211&gt; 32

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; PCR primer

&lt;400&gt; 385

cggcgaattc cacgaaccac tcgcaagttc ag

32

&lt;210&gt; 386

&lt;211&gt; 30

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; PCR primer

&lt;400&gt; 386

cggctcgagt tagcttgggc ctgtgattgc

30

&lt;210&gt; 387

&lt;211&gt; 20

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 387

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Phe | Phe | Lys | Trp | Leu | Leu | Ser | Cys | Cys | Pro | Gly | Ser | Ser | Gln | Ile | Ala |
| 1   |     |     |     | 5   |     |     |     |     | 10  |     |     |     | 15  |     |     |

|     |     |     |     |
|-----|-----|-----|-----|
| Ala | Ala | Ala | Ser |
|     |     |     | 20  |

&lt;210&gt; 388

&lt;211&gt; 19

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 388

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Leu | Ser | Cys | Cys | Pro | Gly | Ser | Ser | Gln | Ile | Ala | Ala | Ala | Ser | Thr | Gln |
| 1   |     |     |     | 5   |     |     |     | 10  |     |     |     |     | 15  |     |     |

|     |     |     |
|-----|-----|-----|
| Pro | Glu | Asp |
|-----|-----|-----|

&lt;210&gt; 389

&lt;211&gt; 20

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 389

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ala | Ala | Ala | Ala | Ser | Thr | Gln | Pro | Glu | Asp | Asp | Ile | Asn | Thr | Gln | Arg |
| 1   |     |     |     | 5   |     |     |     | 10  |     |     |     |     | 15  |     |     |

|     |     |     |     |
|-----|-----|-----|-----|
| Lys | Lys | Ser | Gln |
|     |     |     | 20  |

&lt;210&gt; 390

<211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 390  
 Thr Gln Pro Glu Asp Asp Ile Asn Thr Gln Arg Lys Lys Ser Gln Glu  
 1 5 10 15  
 Lys Met Arg Glu  
 20

<210> 391  
 <211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 391  
 Asp Ile Asn Thr Gln Arg Lys Lys Ser Gln Glu Lys Met Arg Glu Val  
 1 5 10 15  
 Thr Asp Ser Pro  
 20

<210> 392  
 <211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 392  
 Arg Lys Lys Ser Gln Glu Lys Met Arg Glu Val Thr Asp Ser Pro Gly  
 1 5 10 15  
 Arg Pro Arg Glu  
 20

<210> 393  
 <211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 393  
 Glu Lys Met Arg Glu Val Thr Asp Ser Pro Gly Arg Pro Arg Glu Leu  
 1 5 10 15  
 Thr Ile Pro Gln  
 20

<210> 394  
 <211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 394  
 Val Thr Asp Ser Pro Gly Arg Pro Arg Glu Leu Thr Ile Pro Gln Thr

1 5 10 15  
 Ser Ser His Gly  
 20

<210> 395  
 <211> 19  
 <212> PRT  
 <213> Homo sapiens

<400> 395  
 Gly Arg Pro Arg Glu Leu Thr Ile Pro Gln Thr Ser Ser His Gly Ala  
 1 5 10 15  
 Asn Arg Phe

<210> 396  
 <211> 19  
 <212> PRT  
 <213> Homo sapiens

<400> 396  
 Met Asn Lys Leu Tyr Ile Gly Asn Leu Ser Glu Asn Ala Ala Pro Ser  
 1 5 10 15  
 Asp Leu Glu

<210> 397  
 <211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 397  
 Ser Glu Asn Ala Ala Pro Ser Asp Leu Glu Ser Ile Phe Lys Asp Ala  
 1 5 10 15  
 Lys Ile Pro Val  
 20

<210> 398  
 <211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 398  
 Ser Ile Phe Lys Asp Ala Lys Ile Pro Val Ser Gly Pro Phe Leu Val  
 1 5 10 15  
 Lys Thr Gly Tyr  
 20

<210> 399

<211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 399  
 Ser Gly Pro Phe Leu Val Lys Thr Gly Tyr Ala Phe Val Asp Cys Pro  
 1 5 10 15  
 Asp Glu Ser Trp  
 20

<210> 400  
 <211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 400  
 Ala Phe Val Asp Cys Pro Asp Glu Ser Trp Ala Leu Lys Ala Ile Glu  
 1 5 10 15  
 Ala Leu Ser Gly  
 20

<210> 401  
 <211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 401  
 Ala Leu Lys Ala Ile Glu Ala Leu Ser Gly Lys Ile Glu Leu His Gly  
 1 5 10 15  
 Lys Pro Ile Glu  
 20

<210> 402  
 <211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 402  
 Lys Ile Glu Leu His Gly Lys Pro Ile Glu Val Glu His Ser Val Pro  
 1 5 10 15  
 Lys Arg Gln Arg  
 20

<210> 403  
 <211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 403  
 Val Glu His Ser Val Pro Lys Arg Gln Arg Ile Arg Lys Leu Gln Ile

1                    5                    10                    15  
 Arg Asn Ile Pro  
                   20

<210> 404  
 <211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 404  
 Ile Arg Lys Leu Gln Ile Arg Asn Ile Pro Pro His Leu Gln Trp Glu  
   1                  5                  10                  15  
 Val Leu Asp Ser  
                   20

<210> 405  
 <211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 405  
 Ala Val Val Asn Val Thr Tyr Ser Ser Lys Asp Gln Ala Arg Gln Ala  
   1                  5                  10                  15  
 Leu Asp Lys Leu  
                   20

<210> 406  
 <211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 406  
 Asp Gln Ala Arg Gln Ala Leu Asp Lys Leu Asn Gly Phe Gln Leu Glu  
   1                  5                  10                  15  
 Asn Phe Thr Leu  
                   20

<210> 407  
 <211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 407  
 Asn Gly Phe Gln Leu Glu Asn Phe Thr Leu Lys Val Ala Tyr Ile Pro  
   1                  5                  10                  15  
 Asp Glu Thr Ala  
                   20

<210> 408

<400> 412  
Lys Gln Lys Pro Cys Asp Leu Pro Leu Arg Leu Leu Val Pro Thr Gln

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<210> 413
<211> 20
<212> PRT
<213> Homo sapiens
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```
<210> 414
<211> 20
<212> PRT
<213> Homo sapiens
```

```
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<211> 20
<212> PRT
<213> Homo sapiens
```

```
<210> 416
<211> 20
<212> PRT
<213> Homo sapiens
```

<210> 417



<211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 417  
 Ala Glu Lys Ser Ile Thr Ile Leu Ser Thr Pro Glu Gly Thr Ser Ala  
 1 5 10 15  
 Ala Cys Lys Ser  
 20

<210> 418  
 <211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 418  
 Pro Glu Gly Thr Ser Ala Ala Cys Lys Ser Ile Leu Glu Ile Met His  
 1 5 10 15  
 Lys Glu Ala Gln  
 20

<210> 419  
 <211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 419  
 Ile Leu Glu Ile Met His Lys Glu Ala Gln Asp Ile Lys Phe Thr Glu  
 1 5 10 15  
 Glu Ile Pro Leu  
 20

<210> 420  
 <211> 455  
 <212> DNA  
 <213> Homo sapiens

<400> 420  
 gaagacatgc ttacttcccc ttcaccttcc ttcattgatgt gggaagagtg ctgcaaccca 60  
 gccctagcca acgccgcatg agagggagtg tgccgagggc ttctgagaag gtttctctca 120  
 catctagaaa gaagcgctta agatgtggca gcccctcttc ttcaagtggc tcttgctctg 180  
 ttgccctggg agttctcaaa ttgctgcagc agcctccacc cagcctgagg atgacatcaa 240  
 tacacagagg aagaagagtc aggaaaagat gagagaagtt acagactctc ctgggcgacc 300  
 ccgagagctt accattcctc agacttcttc acatggtgct aacagatttg ttcctaaaag 360  
 taaagctcta gaggccgtca aattggcaat agaagccggg ttccaccata ttgattctgc 420  
 acatgtttac aataatgagg agcaggttgg actgg 455

<210> 421  
 <211> 24  
 <212> DNA  
 <213> Artificial Sequence

&lt;220&gt;

&lt;223&gt; PCR primer

&lt;400&gt; 421

actagtgtcc gcgtggcggc ctac

24

&lt;210&gt; 422

&lt;211&gt; 34

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; PCR primer

&lt;400&gt; 422

catgagaatt catcacatgc ccttgaaggc tccc

34

&lt;210&gt; 423

&lt;211&gt; 161

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 423

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Met | Gln | His | His | His | His | His | His | His | Thr | Ser | Val | Arg | Val | Ala | Ala |
| 1   |     |     |     | 5   |     |     |     |     | 10  |     |     |     |     | 15  |     |
| Tyr | Phe | Glu | Asn | Phe | Leu | Ala | Ala | Trp | Arg | Pro | Val | Lys | Ala | Ser | Asp |
|     |     |     | 20  |     |     |     |     | 25  |     |     |     |     | 30  |     |     |
| Gly | Asp | Tyr | Tyr | Thr | Leu | Ala | Val | Pro | Met | Gly | Asp | Val | Pro | Met | Asp |
|     |     | 35  |     |     |     |     | 40  |     |     |     |     | 45  |     |     |     |
| Gly | Ile | Ser | Val | Ala | Asp | Ile | Gly | Ala | Ala | Val | Ser | Ser | Ile | Phe | Asn |
|     |     | 50  |     |     |     | 55  |     |     |     | 60  |     |     |     |     |     |
| Ser | Pro | Glu | Glu | Phe | Leu | Gly | Lys | Ala | Val | Gly | Leu | Ser | Ala | Glu | Ala |
| 65  |     |     |     |     | 70  |     |     |     |     | 75  |     |     |     | 80  |     |
| Leu | Thr | Ile | Gln | Gln | Tyr | Ala | Asp | Val | Leu | Ser | Lys | Ala | Leu | Gly | Lys |
|     |     |     | 85  |     |     |     |     | 90  |     |     |     |     |     | 95  |     |
| Glu | Val | Arg | Asp | Ala | Lys | Ile | Thr | Pro | Glu | Ala | Phe | Glu | Lys | Leu | Gly |
|     |     |     | 100 |     |     |     |     | 105 |     |     |     |     | 110 |     |     |
| Phe | Pro | Ala | Ala | Lys | Glu | Ile | Ala | Asn | Met | Cys | Arg | Phe | Tyr | Glu | Met |
|     |     | 115 |     |     |     |     | 120 |     |     |     |     | 125 |     |     |     |
| Lys | Pro | Asp | Arg | Asp | Val | Asn | Leu | Thr | His | Gln | Leu | Asn | Pro | Lys | Val |
|     | 130 |     |     |     |     | 135 |     |     |     |     | 140 |     |     |     |     |
| Lys | Ser | Phe | Ser | Gln | Phe | Ile | Ser | Glu | Asn | Gln | Gly | Ala | Phe | Lys | Gly |
| 145 |     |     |     | 150 |     |     |     |     | 155 |     |     |     |     | 160 |     |

Met

&lt;210&gt; 424

&lt;211&gt; 489

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 424

```

atgcagcatc accaccatca ccaccacact agtgtccgcg tggcggccta ctttgaaaac 60
tttctcgcgg cgtggcggcc cgtgaaagcc tctgatggag attactacac cttggctgta 120
ccgatgggag atgtaccaat ggatggatc tctgttgctg atattggagc agccgtctct 180
agcattttta attctccaga ggaattttta ggcaaggccg tggggctcag tgcagaagca 240
ctaacaatac agcaatatgc tgatgttttg tccaaggctt tggggaaaga agtccgagat 300
gcaaagatta ccccggaagc ttctcgagaag ctgggattcc ctgcagcaaa ggaaatagcc 360
aatatgtgtc gtttctatga aatgaagcca gaccgagatg tcaatctcac ccaccaacta 420
aatcccaaag tcaaaaagctt cagccagttt atctcagaga accagggagc cttcaagggc 480
atgtgatga 489

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<210> 425

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 425

aacaaactgt atatcggaaa cctcagcgag aa

32

<210> 426

<211> 33

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 426

ccatagaatt cattacttcc gtcttgactg agg

33

<210> 427

<211> 586

<212> PRT

<213> Homo sapiens

<400> 427

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Met Gln His His His His His Asn Lys Leu Tyr Ile Gly Asn Leu
 1          5          10          15
Ser Glu Asn Ala Ala Pro Ser Asp Leu Glu Ser Ile Phe Lys Asp Ala
 20          25          30
Lys Ile Pro Val Ser Gly Pro Phe Leu Val Lys Thr Gly Tyr Ala Phe
 35          40          45
Val Asp Cys Pro Asp Glu Ser Trp Ala Leu Lys Ala Ile Glu Ala Leu
 50          55          60
Ser Gly Lys Ile Glu Leu His Gly Lys Pro Ile Glu Val Glu His Ser
 65          70          75          80
Val Pro Lys Arg Gln Arg Ile Arg Lys Leu Gln Ile Arg Asn Ile Pro
 85          90          95
Pro His Leu Gln Trp Glu Val Leu Asp Ser Leu Leu Val Gln Tyr Gly
100          105          110
Val Val Glu Ser Cys Glu Gln Val Asn Thr Asp Ser Glu Thr Ala Val
115          120          125

```

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Val | Asn | Val | Thr | Tyr | Ser | Ser | Lys | Asp | Gln | Ala | Arg | Gln | Ala | Leu | Asp |
| 130 |     |     |     |     |     | 135 |     |     |     |     | 140 |     |     |     |     |
| Lys | Leu | Asn | Gly | Phe | Gln | Leu | Glu | Asn | Phe | Thr | Leu | Lys | Val | Ala | Tyr |
| 145 |     |     |     |     | 150 |     |     |     |     | 155 |     |     |     |     | 160 |
| Ile | Pro | Asp | Glu | Thr | Ala | Ala | Gln | Gln | Asn | Pro | Leu | Gln | Gln | Pro | Arg |
|     |     |     |     | 165 |     |     |     |     | 170 |     |     |     |     | 175 |     |
| Gly | Arg | Arg | Gly | Leu | Gly | Gln | Arg | Gly | Ser | Ser | Arg | Gln | Gly | Ser | Pro |
|     |     |     | 180 |     |     |     |     | 185 |     |     |     |     | 190 |     |     |
| Gly | Ser | Val | Ser | Lys | Gln | Lys | Pro | Cys | Asp | Leu | Pro | Leu | Arg | Leu | Leu |
|     |     | 195 |     |     |     |     | 200 |     |     |     |     | 205 |     |     |     |
| Val | Pro | Thr | Gln | Phe | Val | Gly | Ala | Ile | Ile | Gly | Lys | Glu | Gly | Ala | Thr |
|     |     | 210 |     |     |     | 215 |     |     |     |     |     | 220 |     |     |     |
| Ile | Arg | Asn | Ile | Thr | Lys | Gln | Thr | Gln | Ser | Lys | Ile | Asp | Val | His | Arg |
| 225 |     |     |     |     | 230 |     |     |     |     | 235 |     |     |     |     | 240 |
| Lys | Glu | Asn | Ala | Gly | Ala | Ala | Glu | Lys | Ser | Ile | Thr | Ile | Leu | Ser | Thr |
|     |     |     |     | 245 |     |     |     |     | 250 |     |     |     |     | 255 |     |
| Pro | Glu | Gly | Thr | Ser | Ala | Ala | Cys | Lys | Ser | Ile | Leu | Glu | Ile | Met | His |
|     |     |     | 260 |     |     |     |     | 265 |     |     |     |     | 270 |     |     |
| Lys | Glu | Ala | Gln | Asp | Ile | Lys | Phe | Thr | Glu | Glu | Ile | Pro | Leu | Lys | Ile |
|     |     | 275 |     |     |     |     | 280 |     |     |     |     | 285 |     |     |     |
| Leu | Ala | His | Asn | Asn | Phe | Val | Gly | Arg | Leu | Ile | Gly | Lys | Glu | Gly | Arg |
|     |     | 290 |     |     |     | 295 |     |     |     |     | 300 |     |     |     |     |
| Asn | Leu | Lys | Lys | Ile | Glu | Gln | Asp | Thr | Asp | Thr | Lys | Ile | Thr | Ile | Ser |
| 305 |     |     |     |     | 310 |     |     |     |     | 315 |     |     |     |     | 320 |
| Pro | Leu | Gln | Glu | Leu | Thr | Leu | Tyr | Asn | Pro | Glu | Arg | Thr | Ile | Thr | Val |
|     |     |     |     | 325 |     |     |     |     | 330 |     |     |     |     | 335 |     |
| Lys | Gly | Asn | Val | Glu | Thr | Cys | Ala | Lys | Ala | Glu | Glu | Glu | Ile | Met | Lys |
|     |     | 340 |     |     |     |     |     | 345 |     |     |     |     | 350 |     |     |
| Lys | Ile | Arg | Glu | Ser | Tyr | Glu | Asn | Asp | Ile | Ala | Ser | Met | Asn | Leu | Gln |
|     |     | 355 |     |     |     |     | 360 |     |     |     |     | 365 |     |     |     |
| Ala | His | Leu | Ile | Pro | Gly | Leu | Asn | Leu | Asn | Ala | Leu | Gly | Leu | Phe | Pro |
|     |     | 370 |     |     |     | 375 |     |     |     |     | 380 |     |     |     |     |
| Pro | Thr | Ser | Gly | Met | Pro | Pro | Pro | Thr | Ser | Gly | Pro | Pro | Ser | Ala | Met |
| 385 |     |     |     |     | 390 |     |     |     |     | 395 |     |     |     |     | 400 |
| Thr | Pro | Pro | Tyr | Pro | Gln | Phe | Glu | Gln | Ser | Glu | Thr | Glu | Thr | Val | His |
|     |     |     |     | 405 |     |     |     |     | 410 |     |     |     |     | 415 |     |
| Leu | Phe | Ile | Pro | Ala | Leu | Ser | Val | Gly | Ala | Ile | Ile | Gly | Lys | Gln | Gly |
|     |     |     | 420 |     |     |     |     | 425 |     |     |     |     | 430 |     |     |
| Gln | His | Ile | Lys | Gln | Leu | Ser | Arg | Phe | Ala | Gly | Ala | Ser | Ile | Lys | Ile |
|     |     | 435 |     |     |     |     | 440 |     |     |     |     | 445 |     |     |     |
| Ala | Pro | Ala | Glu | Ala | Pro | Asp | Ala | Lys | Val | Arg | Met | Val | Ile | Ile | Thr |
|     |     | 450 |     |     |     | 455 |     |     |     |     | 460 |     |     |     |     |
| Gly | Pro | Pro | Glu | Ala | Gln | Phe | Lys | Ala | Gln | Gly | Arg | Ile | Tyr | Gly | Lys |
| 465 |     |     |     |     | 470 |     |     |     |     | 475 |     |     |     |     | 480 |
| Ile | Lys | Glu | Glu | Asn | Phe | Val | Ser | Pro | Lys | Glu | Glu | Val | Lys | Leu | Glu |
|     |     |     |     | 485 |     |     |     |     | 490 |     |     |     |     | 495 |     |
| Ala | His | Ile | Arg | Val | Pro | Ser | Phe | Ala | Ala | Gly | Arg | Val | Ile | Gly | Lys |
|     |     |     | 500 |     |     |     |     | 505 |     |     |     |     | 510 |     |     |
| Gly | Gly | Lys | Thr | Val | Asn | Glu | Leu | Gln | Asn | Leu | Ser | Ser | Ala | Glu | Val |
|     |     | 515 |     |     |     |     | 520 |     |     |     |     | 525 |     |     |     |
| Val | Val | Pro | Arg | Asp | Gln | Thr | Pro | Asp | Glu | Asn | Asp | Gln | Val | Val | Val |
|     |     | 530 |     |     |     | 535 |     |     |     |     | 540 |     |     |     |     |
| Lys | Ile | Thr | Gly | His | Phe | Tyr | Ala | Cys | Gln | Val | Ala | Gln | Arg | Lys | Ile |
| 545 |     |     |     |     | 550 |     |     |     |     | 555 |     |     |     |     | 560 |

130 135 140 145 150 155 160 165 170 175 180 185 190 195 200 205 210 215 220 225 230 235 240 245 250 255 260 265 270 275 280 285 290 295 300 305 310 315 320 325 330 335 340 345 350 355 360 365 370 375 380 385 390 395 400 405 410 415 420 425 430 435 440 445 450 455 460 465 470 475 480 485 490 495 500 505 510 515 520 525 530 535 540 545 550 555 560

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<210> 428
<211> 1764
<212> DNA
<213> Homo sapiens
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<210> 429
<211> 35
<212> DNA
<213> Artificial Sequence
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<400> 429
ccatggaatt cattatttca atataagata atctc
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<210> 430  
 <211> 881  
 <212> PRT  
 <213> Homo sapiens

<400> 430

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Met | Gln | His | His | His | His | His | His | Gly | Val | Gln | Leu | Gln | Asp | Asn | Gly |
| 1   |     |     |     | 5   |     |     |     |     | 10  |     |     |     |     | 15  |     |
| Tyr | Asn | Gly | Leu | Leu | Ile | Ala | Ile | Asn | Pro | Gln | Val | Pro | Glu | Asn | Gln |
|     |     |     | 20  |     |     |     |     | 25  |     |     |     |     | 30  |     |     |
| Asn | Leu | Ile | Ser | Asn | Ile | Lys | Glu | Met | Ile | Thr | Glu | Ala | Ser | Phe | Tyr |
|     |     | 35  |     |     |     |     | 40  |     |     |     |     | 45  |     |     |     |
| Leu | Phe | Asn | Ala | Thr | Lys | Arg | Val | Phe | Phe | Arg | Asn | Ile | Lys | Ile |     |
|     | 50  |     |     |     |     | 55  |     |     |     | 60  |     |     |     |     |     |
| Leu | Ile | Pro | Ala | Thr | Trp | Lys | Ala | Asn | Asn | Asn | Ser | Lys | Ile | Lys | Gln |
| 65  |     |     |     |     | 70  |     |     |     |     | 75  |     |     |     |     | 80  |
| Glu | Ser | Tyr | Glu | Lys | Ala | Asn | Val | Ile | Val | Thr | Asp | Trp | Tyr | Gly | Ala |
|     |     |     |     | 85  |     |     |     |     | 90  |     |     |     |     | 95  |     |
| His | Gly | Asp | Asp | Pro | Tyr | Thr | Leu | Gln | Tyr | Arg | Gly | Cys | Gly | Lys | Glu |
|     |     |     | 100 |     |     |     |     | 105 |     |     |     |     | 110 |     |     |
| Gly | Lys | Tyr | Ile | His | Phe | Thr | Pro | Asn | Phe | Leu | Leu | Asn | Asp | Asn | Leu |
|     |     | 115 |     |     |     |     | 120 |     |     |     |     | 125 |     |     |     |
| Thr | Ala | Gly | Tyr | Gly | Ser | Arg | Gly | Arg | Val | Phe | Val | His | Glu | Trp | Ala |
|     | 130 |     |     |     |     | 135 |     |     |     |     | 140 |     |     |     |     |
| His | Leu | Arg | Trp | Gly | Val | Phe | Asp | Glu | Tyr | Asn | Asn | Asp | Lys | Pro | Phe |
| 145 |     |     |     |     | 150 |     |     |     |     | 155 |     |     |     |     | 160 |
| Tyr | Ile | Asn | Gly | Gln | Asn | Gln | Ile | Lys | Val | Thr | Arg | Cys | Ser | Ser | Asp |
|     |     |     | 165 |     |     |     |     |     | 170 |     |     |     |     | 175 |     |
| Ile | Thr | Gly | Ile | Phe | Val | Cys | Glu | Lys | Gly | Pro | Cys | Pro | Gln | Glu | Asn |
|     |     | 180 |     |     |     |     |     | 185 |     |     |     |     | 190 |     |     |
| Cys | Ile | Ile | Ser | Lys | Leu | Phe | Lys | Glu | Gly | Cys | Thr | Phe | Ile | Tyr | Asn |
|     | 195 |     |     |     |     |     | 200 |     |     |     |     | 205 |     |     |     |
| Ser | Thr | Gln | Asn | Ala | Thr | Ala | Ser | Ile | Met | Phe | Met | Gln | Ser | Leu | Ser |
|     | 210 |     |     |     |     | 215 |     |     |     |     | 220 |     |     |     |     |
| Ser | Val | Val | Glu | Phe | Cys | Asn | Ala | Ser | Thr | His | Asn | Gln | Glu | Ala | Pro |
| 225 |     |     |     |     | 230 |     |     |     |     | 235 |     |     |     |     | 240 |
| Asn | Leu | Gln | Asn | Gln | Met | Cys | Ser | Leu | Arg | Ser | Ala | Trp | Asp | Val | Ile |
|     |     |     | 245 |     |     |     |     |     | 250 |     |     |     |     | 255 |     |
| Thr | Asp | Ser | Ala | Asp | Phe | His | His | Ser | Phe | Pro | Met | Asn | Gly | Thr | Glu |
|     |     | 260 |     |     |     |     |     | 265 |     |     |     |     | 270 |     |     |
| Leu | Pro | Pro | Pro | Pro | Thr | Phe | Ser | Leu | Val | Glu | Ala | Gly | Asp | Lys | Val |
|     | 275 |     |     |     |     |     | 280 |     |     |     |     | 285 |     |     |     |
| Val | Cys | Leu | Val | Leu | Asp | Val | Ser | Ser | Lys | Met | Ala | Glu | Ala | Asp | Arg |
|     | 290 |     |     |     |     | 295 |     |     |     |     | 300 |     |     |     |     |
| Leu | Leu | Gln | Leu | Gln | Gln | Ala | Ala | Glu | Phe | Tyr | Leu | Met | Gln | Ile | Val |
| 305 |     |     |     |     | 310 |     |     |     |     | 315 |     |     |     |     | 320 |
| Glu | Ile | His | Thr | Phe | Val | Gly | Ile | Ala | Ser | Phe | Asp | Ser | Lys | Gly | Glu |
|     |     |     |     | 325 |     |     |     |     | 330 |     |     |     |     | 335 |     |
| Ile | Arg | Ala | Gln | Leu | His | Gln | Ile | Asn | Ser | Asn | Asp | Asp | Arg | Lys | Leu |
|     |     | 340 |     |     |     |     |     | 345 |     |     |     |     | 350 |     |     |
| Leu | Val | Ser | Tyr | Leu | Pro | Thr | Thr | Val | Ser | Ala | Lys | Thr | Asp | Ile | Ser |
|     | 355 |     |     |     |     |     | 360 |     |     |     |     | 365 |     |     |     |
| Ile | Cys | Ser | Gly | Leu | Lys | Lys | Gly | Phe | Glu | Val | Val | Glu | Lys | Leu | Asn |
|     | 370 |     |     |     |     | 375 |     |     |     |     |     | 380 |     |     |     |

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Gly Lys Ala Tyr Gly Ser Val Met Ile Leu Val Thr Ser Gly Asp Asp  
 385 390 395 400  
 Lys Leu Leu Gly Asn Cys Leu Pro Thr Val Leu Ser Ser Gly Ser Thr  
 405 410 415  
 Ile His Ser Ile Ala Leu Gly Ser Ser Ala Ala Pro Asn Leu Glu Glu  
 420 425 430  
 Leu Ser Arg Leu Thr Gly Gly Leu Lys Phe Phe Val Pro Asp Ile Ser  
 435 440 445  
 Asn Ser Asn Ser Met Ile Asp Ala Phe Ser Arg Ile Ser Ser Gly Thr  
 450 455 460  
 Gly Asp Ile Phe Gln Gln His Ile Gln Leu Glu Ser Thr Gly Glu Asn  
 465 470 475 480  
 Val Lys Pro His His Gln Leu Lys Asn Thr Val Thr Val Asp Asn Thr  
 485 490 495  
 Val Gly Asn Asp Thr Met Phe Leu Val Thr Trp Gln Ala Ser Gly Pro  
 500 505 510  
 Pro Glu Ile Ile Leu Phe Asp Pro Asp Gly Arg Lys Tyr Tyr Thr Asn  
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 Asn Phe Ile Thr Asn Leu Thr Phe Arg Thr Ala Ser Leu Trp Ile Pro  
 530 535 540  
 Gly Thr Ala Lys Pro Gly His Trp Thr Tyr Thr Leu Asn Asn Thr His  
 545 550 555 560  
 His Ser Leu Gln Ala Leu Lys Val Thr Val Thr Ser Arg Ala Ser Asn  
 565 570 575  
 Ser Ala Val Pro Pro Ala Thr Val Glu Ala Phe Val Glu Arg Asp Ser  
 580 585 590  
 Leu His Phe Pro His Pro Val Met Ile Tyr Ala Asn Val Lys Gln Gly  
 595 600 605  
 Phe Tyr Pro Ile Leu Asn Ala Thr Val Thr Ala Thr Val Glu Pro Glu  
 610 615 620  
 Thr Gly Asp Pro Val Thr Leu Arg Leu Leu Asp Asp Gly Ala Gly Ala  
 625 630 635 640  
 Asp Val Ile Lys Asn Asp Gly Ile Tyr Ser Arg Tyr Phe Phe Ser Phe  
 645 650 655  
 Ala Ala Asn Gly Arg Tyr Ser Leu Lys Val His Val Asn His Ser Pro  
 660 665 670  
 Ser Ile Ser Thr Pro Ala His Ser Ile Pro Gly Ser His Ala Met Tyr  
 675 680 685  
 Val Pro Gly Tyr Thr Ala Asn Gly Asn Ile Gln Met Asn Ala Pro Arg  
 690 695 700  
 Lys Ser Val Gly Arg Asn Glu Glu Glu Arg Lys Trp Gly Phe Ser Arg  
 705 710 715 720  
 Val Ser Ser Gly Gly Ser Phe Ser Val Leu Gly Val Pro Ala Gly Pro  
 725 730 735  
 His Pro Asp Val Phe Pro Pro Cys Lys Ile Ile Asp Leu Glu Ala Val  
 740 745 750  
 Lys Val Glu Glu Glu Leu Thr Leu Ser Trp Thr Ala Pro Gly Glu Asp  
 755 760 765  
 Phe Asp Gln Gly Gln Ala Thr Ser Tyr Glu Ile Arg Met Ser Lys Ser  
 770 775 780  
 Leu Gln Asn Ile Gln Asp Asp Phe Asn Asn Ala Ile Leu Val Asn Thr  
 785 790 795 800  
 Ser Lys Arg Asn Pro Gln Gln Ala Gly Ile Arg Glu Ile Phe Thr Phe  
 805 810 815

Ser Pro Gln Ile Ser Thr Asn Gly Pro Glu His Gln Pro Asn Gly Glu  
                   820                  825                  830  
 Thr His Glu Ser His Arg Ile Tyr Val Ala Ile Arg Ala Met Asp Arg  
                   835                  840                  845  
 Asn Ser Leu Gln Ser Ala Val Ser Asn Ile Ala Gln Ala Pro Leu Phe  
                   850                  855                  860  
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 Lys

<210> 431  
 <211> 2646  
 <212> DNA  
 <213> Homo sapiens

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 ctcatcgcaa ttaatcctca ggtacctgag aatcagaacc tcatctcaaa cattaaggaa 120  
 atgataactg aagcttcatt ttacctatit aatgctacca agagaagagt atttttcaga 180  
 aatataaaga ttttaatacc tgccacatgg aaagctaata ataacagcaa aataaaacaa 240  
 gaatcatatg aaaaggcaaa tgtcatagtg actgactggg atgggggcaca tggagatgat 300  
 ccatacaccc tacaatacac aggggtgtgga aaagagggaa aatacattca tttcacacct 360  
 aatttcctac tgaatgataa cttaacagct ggctacggat cagcaggccg agtggtttgtc 420  
 catgaatggg cccacctccg ttgggggtgtg ttcatgagat ataacaatga caaacctttc 480  
 tacataaatg ggcaaaatca aattaaagtg acaaggtgtt catctgacat cacaggcatt 540  
 tttgtgtgtg aaaaagggtcc ttgcccccaa gaaaactgta ttattagtaa gcttttttaa 600  
 gaaggatgca cctttatcta caatagcacc caaaatgcaa ctgcatcaat aatgttcattg 660  
 caaagtttat cttctgtggg tgaattttgt aatgcaagta cccacaacca agaagcacca 720  
 aacctacaga accagatgtg cagcctcaga agtgcatggg atgtaatcac agactctgct 780  
 gactttcacc acagctttcc catgaacggg actgagcttc cacctcctcc cacattctcg 840  
 cttgtagagg ctgggtgaca agtggtctgt tttagtctgg atgtgtccag caagatggca 900  
 gaggtgaca gactccttca actacaacaa gccgcagaat tttatttgat gcagattgtt 960  
 gaaattcata ctttcgtggg cattgccagt ttgcagacga aaggagagat cagagccag 1020  
 ctacaccaa ttaacagcaa tgatgatcga aagttgctgg tttcatatct gccaccact 1080  
 gtatcagcta aaacagacat cagcatttgt tcagggtcta agaaaggatt tgaggtgggt 1140  
 gaaaaactga atggaaaagc ttatggctct gtgatgatat tagtgaccag cggagatgat 1200  
 aagcttcttg gcaattgctt acccactgtg ctacagcagt gttcaacaat tcaactccatt 1260  
 gccctgggtt catctgcagc cccaaatctg gaggaattat cacgtcttac aggaggttta 1320  
 aagttctttg ttccagatat atcaaaactcc aatagcatga ttgatgcttt cagtagaatt 1380  
 tcctctggaa ctggagacat ttccagcaa catattcagc ttgaaagtac aggtgaaaat 1440  
 gtcaaacctc accatcaatt gaaaaacaca gtgactgtgg ataatactgt gggcaacgac 1500  
 actatgtttc tagttacgtg gcaggccagt ggtcctcctg agattatatt atttgatcct 1560  
 gatggacgaa aatactacac aaataatttt atcaccaatc taacttttgc gacagctagt 1620  
 ctttggtatc caggaacagc taagcctggg cactggactt acacctgaa caatacccat 1680  
 cattctctgc aagccctgaa agtgacagt acctctcgcg cctccaactc agctgtgcc 1740  
 ccagccactg tggagccctt tgtggaaaga gacagcctcc attttcctca tcctgtgatg 1800  
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 gttgagccag agactggaga tcctgttacg ctgagactcc ttgatgatgg agcaggtgct 1920  
 gatgttataa aaaatgatgg aatttactcg aggtattttt tctcctttgc tgcaaattgg 1980  
 agatatagct tgaaagtgca tgtcaatcac tctcccagca taagcaccac agcccactct 2040  
 attccaggga gtcatgctat gtatgtacca ggttacacag caaacggtaa tattcagatg 2100  
 aatgctccaa ggaaatcagt aggcagaaat gaggaggagc gaaagtgggg ctttagccga 2160



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gtcagctcag gaggtcctt ttcagtgtg ggagttccag ctggcccca cctgatgtg 2220
tttccaccat gcaaaattat tgacctggaa gctgtaaaag tagaagagga attgacccta 2280
tcttggacag cacctggaga agactttgat cagggccagg ctacaagcta tgaaataaga 2340
atgagtaaaa gtctacagaa tatccaagat gactttaaca atgctatttt agtaaataca 2400
tcaaagcgaa atcctcagca agctggcatc agggagatat ttacgttctc accccaaatt 2460
tccacgaatg gacctgaaca tcagccaaat ggagaaacac atgaaagcca cagaatttat 2520
gttgcaatac gagcaatgga taggaactcc ttacagtctg ctgtatctaa cattgcccag 2580
gcgcctctgt ttattccccc caattctgat cctgtacctg ccagagatta tcttatattg 2640
aaataa 2646

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<210> 432

<211> 36

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 432

cgctgctcg agtcattaat attcatcaga aaatgg

36

<210> 433

<211> 371

<212> PRT

<213> Homo sapiens

<400> 433

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Met Gln His His His His His Trp Gln Pro Leu Phe Phe Lys Trp
 1          5          10          15
Leu Leu Ser Cys Cys Pro Gly Ser Ser Gln Ile Ala Ala Ala Ala Ser
          20          25          30
Thr Gln Pro Glu Asp Asp Ile Asn Thr Gln Arg Lys Lys Ser Gln Glu
          35          40          45
Lys Met Arg Glu Val Thr Asp Ser Pro Gly Arg Pro Arg Glu Leu Thr
          50          55          60
Ile Pro Gln Thr Ser Ser His Gly Ala Asn Arg Phe Val Pro Lys Ser
65          70          75          80
Lys Ala Leu Glu Ala Val Lys Leu Ala Ile Glu Ala Gly Phe His His
          85          90          95
Ile Asp Ser Ala His Val Tyr Asn Asn Glu Glu Gln Val Gly Leu Ala
          100          105          110
Ile Arg Ser Lys Ile Ala Asp Gly Ser Val Lys Arg Glu Asp Ile Phe
          115          120          125
Tyr Thr Ser Lys Leu Trp Ser Asn Ser His Arg Pro Glu Leu Val Arg
          130          135          140
Pro Ala Leu Glu Arg Ser Leu Lys Asn Leu Gln Leu Asp Tyr Val Asp
145          150          155          160
Leu Tyr Leu Ile His Phe Pro Val Ser Val Lys Pro Gly Glu Glu Val
          165          170          175
Ile Pro Lys Asp Glu Asn Gly Lys Ile Leu Phe Asp Thr Val Asp Leu
          180          185          190
Cys Ala Thr Trp Glu Ala Met Glu Lys Cys Lys Asp Ala Gly Leu Ala
          195          200          205
Lys Ser Ile Gly Val Ser Asn Phe Asn His Arg Leu Leu Glu Met Ile

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|                     |                         |                     |
|---------------------|-------------------------|---------------------|
| 210                 | 215                     | 220                 |
| Leu Asn Lys Pro Gly | Leu Lys Tyr Lys Pro Val | Cys Asn Gln Val Glu |
| 225                 | 230                     | 235                 |
| Cys His Pro Tyr Phe | Asn Gln Arg Lys Leu Leu | Asp Phe Cys Lys Ser |
| 245                 | 250                     | 255                 |
| Lys Asp Ile Val Leu | Val Ala Tyr Ser Ala Leu | Gly Ser His Arg Glu |
| 260                 | 265                     | 270                 |
| Glu Pro Trp Val Asp | Pro Asn Ser Pro Val Leu | Leu Glu Asp Pro Val |
| 275                 | 280                     | 285                 |
| Leu Cys Ala Leu Ala | Lys Lys His Lys Arg Thr | Pro Ala Leu Ile Ala |
| 290                 | 295                     | 300                 |
| Leu Arg Tyr Gln Leu | Gln Arg Gly Val Val Val | Leu Ala Lys Ser Tyr |
| 305                 | 310                     | 315                 |
| Asn Glu Gln Arg Ile | Arg Gln Asn Val Gln Val | Phe Glu Phe Gln Leu |
| 325                 | 330                     | 335                 |
| Thr Ser Glu Glu Met | Lys Ala Ile Asp Gly Leu | Asn Arg Asn Val Arg |
| 340                 | 345                     | 350                 |
| Tyr Leu Thr Leu Asp | Ile Phe Ala Gly Pro Pro | Asn Tyr Pro Phe Ser |
| 355                 | 360                     | 365                 |
| Asp Glu Tyr         |                         |                     |
| 370                 |                         |                     |

<210> 434  
 <211> 1119  
 <212> DNA  
 <213> Homo sapiens

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| <400> 434  |      |
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| tgccctggga gttctcaaat tgctgcagca gcctccaccc agcctgagga tgacatcaat  | 120  |
| acacagagga agaagagtca ggaaaagatg agagaagtta cagactctcc tgggcgaccc  | 180  |
| cgagagctta ccattcctca gacttcttca catggtgcta acagatttgt tcctaaaagt  | 240  |
| aaagctctag aggccgtcaa attggcaata gaagccgggt tccaccatat tgattctgca  | 300  |
| catgtttaca ataatgagga gcaggttgga ctggccatcc gaagcaagat tgcagatggc  | 360  |
| agtgtgaaga gagaagacat attctacact tcaaagcttt ggagcaattc ccacgacca   | 420  |
| gagttgggtcc gaccagcctt ggaaagggtc ctgaaaaatc ttcaattgga ctatgttgac | 480  |
| ctctatctta ttcatcttcc agtgtctgta aagccagggt aggaagtgat cccaaaagat  | 540  |
| gaaaatggaa aaatactatt tgacacagtg gatctctgtg ccacatggga ggccatggag  | 600  |
| aagtgtaaag atgcaggatt ggccaagtcc atcggggtgt ccaacttcaa ccacaggctg  | 660  |
| ctggagatga tcctcaacaa gccagggtc aagtacaagc ctgtctgcaa ccagggtggaa  | 720  |
| tgtcatcctt acttcaacca gagaaaactg ctggatttct gcaagtcaaa agacattgtt  | 780  |
| ctggttgcct atagtgtctt gggatcccat cgagaagaac catgggtgga cccgaactcc  | 840  |
| ccggtgctct tggaggacct agtcctttgt gccttggcaa aaaagcacia gogaacccca  | 900  |
| gccctgattg cctgcgcta ccagctgcag cgtgggggtg tggtcctggc caagagctac   | 960  |
| aatgagcagc gcatcagaca gaacgtgcag gtgtttgaat tccagttgac ttcagaggag  | 1020 |
| atgaaagcca tagatggcct aaacagaaat gtgcgatatt tgacccttga tatttttgc   | 1080 |
| ggcccccta attatccatt ttctgatgaa tattaatga                          | 1119 |

<210> 435  
 <211> 36  
 <212> DNA  
 <213> Artificial Sequence

<220>  
<223> Primer

<400> 435  
ggatccgccc ccaccatgac atccattcga gctgta 36

<210> 436  
<211> 27  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Primer

<400> 436  
gtcgactcag ctggaccaca gccgcag 27

<210> 437  
<211> 37  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Primer

<400> 437  
ggatccgccc ccaccatgga ctcttgacc ttctgct 37

<210> 438  
<211> 27  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Primer

<400> 438  
gtcgactcag aaatcctttc tcttgac 27

<210> 439  
<211> 933  
<212> DNA  
<213> Homo sapiens

<400> 439  
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gctggagtta tccagtcacc ccggcacgag gtgacagaga tgggacaaga agtgactctg 120  
agatgtaaac caatttcagg acacgactac cttttctggt acagacagac catgatgcgg 180  
ggactggagt tgctcattta ctttaacaac aacgttccga tagatgattc agggatgccc 240  
gaggatcgat tctcagctaa gatgcctaata gcatcattct ccactctgaa gatccagccc 300  
tcagaacca gggactcagc tgtgtacttc tgtgccagca gtttagttgg agcaaact 360  
gaagctttct ttggacaagg caccagactc acagttgtag aggacctgaa caaggtgttc 420  
ccaccgagg tcgctgtgtt tgagccatca gaagcagaga tctcccacac ccaaaaggcc 480  
aactggtgt gctggccac aggtttcttc cctgaccacg tggagctgag ctggtgggtg 540

|            |             |            |            |             |            |     |
|------------|-------------|------------|------------|-------------|------------|-----|
| aatgggaagg | aggtgcacag  | tggggtcagc | acggacccgc | agccccctcaa | ggagcagccc | 600 |
| gccctcaatg | actccagata  | ctgcctgagc | agccgcctga | gggtctcggc  | caccttctgg | 660 |
| cagaaccccc | gcaaccactt  | ccgctgtcaa | gtccagttct | acgggctctc  | ggagaatgac | 720 |
| gagtggaccc | aggatagggc  | caaaccgcgc | accagatcgc | tcagcgccga  | ggcctggggg | 780 |
| agagcagact | gtggctttac  | ctcgggtgtc | taccagcaag | gggtcctgtc  | tgccaccatc | 840 |
| ctctatgaga | tcttgctagg  | gaaggccacc | ctgtatgctg | tgctggtcag  | cgcccttgtg | 900 |
| ttgatggcca | tgggtcaagag | aaaggatttc | tga        |             |            | 933 |

&lt;210&gt; 440

&lt;211&gt; 822

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 440

|            |             |            |             |            |            |     |
|------------|-------------|------------|-------------|------------|------------|-----|
| atgacatcca | ttcgagctgt  | atztatattc | ctgtggctgc  | agctggactt | ggtgaatgga | 60  |
| gagaatgtgg | agcagcatcc  | ttcaaccctg | agtgtccagg  | agggagacag | cgctgttata | 120 |
| aagtgtactt | attcagacag  | tgcctcaaac | tacttccctt  | ggtataagca | agaacttgga | 180 |
| aaaagacctc | agcttattat  | agacattcgt | tcaaagtgtg  | gcgaaaagaa | agaccaacga | 240 |
| attgctgtta | cattgaacaa  | gacagccaaa | catttctccc  | tgacatcac  | agagacccaa | 300 |
| cctgaagact | cggtctgtct  | cttctgtgca | gcaagtatac  | tgaacaccgg | taaccagttc | 360 |
| tattttggga | cagggacaag  | tttgacggtc | attccaaata  | tccagaacct | tgaccctgcc | 420 |
| gtgtaccagc | tgagagactc  | taaatccagt | gacaagtctg  | tctgcctatt | caccgatttt | 480 |
| gattctcaaa | caaagtgtgc  | acaaagtaag | gattctgatg  | tgtatatcac | agacaaaact | 540 |
| gtgctagaca | tgagggtctat | ggacttcaag | agcaacagtg  | ctgtggcctg | gagcaacaaa | 600 |
| tctgactttg | catgtgcaaa  | cgccttcaac | aacagcatta  | ttccagaaga | caccttcttc | 660 |
| cccagcccag | aaagttcctg  | tgatgtcaag | ctggctcgaga | aaagctttga | aacagatacg | 720 |
| aacctaaact | ttcaaaacct  | gtcagtgatt | gggttccgaa  | tcctcctcct | gaaagtggcc | 780 |
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&lt;210&gt; 441

&lt;211&gt; 2311

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 441

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| aagagttggg  | gtttgtctag | gaagagattt | aagcatgctt | gcttaccag  | actcagagaa | 120  |
| gtctccctgt  | tctgtcctag | ctatgttcct | gtgttgtgtg | cattcgtctt | ttccagagca | 180  |
| aaccgcccag  | agtagaagat | ggattggggc | acgtgcaga  | cgatcctggg | gggtgtgaac | 240  |
| aaacactcca  | ccagcattgg | aaagatctgg | ctcaccgtcc | tcttcatttt | tcgcattatg | 300  |
| atcctcgttg  | tggttgcaaa | ggaggtgtgg | ggagatgagc | aggccgactt | tgtctgcaac | 360  |
| accctgcagc  | caggctgcaa | gaacgtgtgc | tacgatcact | acttcccat  | ctccacatc  | 420  |
| cggctatggg  | ccctgcagct | gatcttcgtg | tccagcccag | cgctcctagt | ggccatgcac | 480  |
| gtggcctacc  | ggagacatga | gaagaagagg | aagttcatca | agggggagat | aaagagtga  | 540  |
| tttaaggaca  | tcgaggagat | caaaaccag  | aaggctccga | tcgaaggctc | cctgtgggtg | 600  |
| acctacacaa  | gcagcatctt | cttcgggtgc | atcttcgaag | ccgccttcac | gtacgtcttc | 660  |
| tatgtcatgt  | acgacggctt | ctccatgcag | cggctgggtg | agtgcacgc  | ctggccttgt | 720  |
| cccaacactg  | tggactgctt | tgtgtcccgg | cccacggaga | agactgtctt | cacagtgttc | 780  |
| atgattgcag  | tgtctggaat | ttgcatacct | ctgaatgtca | ctgaattgtg | ttatttgcta | 840  |
| attagatatt  | gttctgggaa | gtcaaaaaag | ccagtttaac | gcattgccc  | gttgttagat | 900  |
| taagaaatag  | acagcatgag | agggatgagg | caaccgcgtc | tcagctgtca | aggctcagtc | 960  |
| gccagcattt  | cccaacacaa | agattctgac | cttaaatgca | accatttgaa | accctgttag | 1020 |
| gcctcaggtg  | aaactccaga | tgccacaatg | agctctgctc | ccctaaagcc | tcaaaacaaa | 1080 |

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<211> 226

<212> PRT

<213> Homo sapiens

<400> 442

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Ile Met Ile Leu Val Val Ala Ala Lys Glu Val Trp Gly Asp Glu Gln  
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Ala Asp Phe Val Cys Asn Thr Leu Gln Pro Gly Cys Lys Asn Val Cys  
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Tyr Asp His Tyr Phe Pro Ile Ser His Ile Arg Leu Trp Ala Leu Gln  
65 70 75 80

Leu Ile Phe Val Ser Ser Pro Ala Leu Leu Val Ala Met His Val Ala  
85 90 95

Tyr Arg Arg His Glu Lys Lys Arg Lys Phe Ile Lys Gly Glu Ile Lys  
100 105 110

Ser Glu Phe Lys Asp Ile Glu Glu Ile Lys Thr Gln Lys Val Arg Ile  
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Glu Gly Ser Leu Trp Trp Thr Tyr Thr Ser Ser Ile Phe Phe Arg Val  
130 135 140

Ile Phe Glu Ala Ala Phe Met Tyr Val Phe Tyr Val Met Tyr Asp Gly  
 145 150 155 160

Phe Ser Met Gln Arg Leu Val Lys Cys Asn Ala Trp Pro Cys Pro Asn  
 165 170 175

Thr Val Asp Cys Phe Val Ser Arg Pro Thr Glu Lys Thr Val Phe Thr  
 180 185 190

Val Phe Met Ile Ala Val Ser Gly Ile Cys Ile Leu Leu Asn Val Thr  
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Pro Val  
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30

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Gly Lys Pro Ile Glu Val Glu His Ser Val Pro Lys Arg Gln Arg Ile  
65 70 75 80  
Arg Lys Leu Gln Ile Arg Asn Ile Pro Pro His Leu Gln Trp Glu Val  
85 90 95  
Leu Asp Ser Leu Leu Val Gln Tyr Gly Val Val Glu Ser Cys Glu Gln  
100 105 110  
Val Asn Thr Asp Ser Glu Thr Ala Val Val Asn Val Thr Tyr Ser Ser  
115 120 125  
Lys Asp Gln Ala Arg Gln Ala Leu Asp Lys Leu Asn Gly Phe Gln Leu  
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Glu Asn Phe Thr Leu Lys Val Ala Tyr Ile Pro Asp Glu Thr Ala Ala  
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Gln Gln Asn Pro Leu Gln Gln Pro Arg Gly Arg Arg Gly Leu Gly Gln  
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180 185 190  
Pro Cys Asp Leu Pro Leu Arg Leu Leu Val Pro Thr Gln Phe Val Gly  
195 200 205  
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210 215 220  
Thr Gln Ser Lys Ile Asp Val His Arg Lys Glu Asn Ala Gly Ala Ala  
225 230 235 240  
Glu Lys Ser Ile Thr Ile Leu Ser Thr Pro Glu Gly Thr Ser Ala Ala

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
|     |     |     |     | 245 |     |     |     |     |     | 250 |     |     |     |     | 255 |
| Cys | Lys | Ser | Ile | Leu | Glu | Ile | Met | His | Lys | Glu | Ala | Gln | Asp | Ile | Lys |
|     |     |     | 260 |     |     |     |     | 265 |     |     |     |     | 270 |     |     |
| Phe | Thr | Glu | Glu | Ile | Pro | Leu | Lys | Ile | Leu | Ala | His | Asn | Asn | Phe | Val |
|     |     | 275 |     |     |     |     | 280 |     |     |     |     | 285 |     |     |     |
| Gly | Arg | Leu | Ile | Gly | Lys | Glu | Gly | Arg | Asn | Leu | Lys | Lys | Ile | Glu | Gln |
|     | 290 |     |     |     |     | 295 |     |     |     |     | 300 |     |     |     |     |
| Asp | Thr | Asp | Thr | Lys | Ile | Thr | Ile | Ser | Pro | Leu | Gln | Glu | Leu | Thr | Leu |
| 305 |     |     |     |     | 310 |     |     |     |     | 315 |     |     |     |     | 320 |
| Tyr | Asn | Pro | Glu | Arg | Thr | Ile | Thr | Val | Lys | Gly | Asn | Val | Glu | Thr | Cys |
|     |     |     |     | 325 |     |     |     |     | 330 |     |     |     |     | 335 |     |
| Ala | Lys | Ala | Glu | Glu | Glu | Ile | Met | Lys | Lys | Ile | Arg | Glu | Ser | Tyr | Glu |
|     |     |     | 340 |     |     |     |     | 345 |     |     |     |     | 350 |     |     |
| Asn | Asp | Ile | Ala | Ser | Met | Asn | Leu | Gln | Ala | His | Leu | Ile | Pro | Gly | Leu |
|     |     | 355 |     |     |     |     | 360 |     |     |     |     | 365 |     |     |     |
| Asn | Leu | Asn | Ala | Leu | Gly | Leu | Phe | Pro | Pro | Thr | Ser | Gly | Met | Pro | Pro |
|     | 370 |     |     |     |     | 375 |     |     |     |     | 380 |     |     |     |     |
| Pro | Thr | Ser | Gly | Pro | Pro | Ser | Ala | Met | Thr | Pro | Pro | Tyr | Pro | Gln | Phe |
| 385 |     |     |     |     | 390 |     |     |     |     | 395 |     |     |     |     | 400 |
| Glu | Gln | Ser | Glu | Thr | Glu | Thr | Val | His | Leu | Phe | Ile | Pro | Ala | Leu | Ser |
|     |     |     |     | 405 |     |     |     |     | 410 |     |     |     |     | 415 |     |
| Val | Gly | Ala | Ile | Ile | Gly | Lys | Gln | Gly | Gln | His | Ile | Lys | Gln | Leu | Ser |
|     |     |     | 420 |     |     |     | 425 |     |     |     |     |     | 430 |     |     |
| Arg | Phe | Ala | Gly | Ala | Ser | Ile | Lys | Ile | Ala | Pro | Ala | Glu | Ala | Pro | Asp |
|     |     | 435 |     |     |     |     | 440 |     |     |     |     | 445 |     |     |     |
| Ala | Lys | Val | Arg | Met | Val | Ile | Ile | Thr | Gly | Pro | Pro | Glu | Ala | Gln | Phe |
|     | 450 |     |     |     |     | 455 |     |     |     | 460 |     |     |     |     |     |
| Lys | Ala | Gln | Gly | Arg | Ile | Tyr | Gly | Lys | Ile | Lys | Glu | Glu | Asn | Phe | Val |
| 465 |     |     |     |     | 470 |     |     |     | 475 |     |     |     |     | 480 |     |
| Ser | Pro | Lys | Glu | Glu | Val | Lys | Leu | Glu | Ala | His | Ile | Arg | Val | Pro | Ser |
|     |     |     |     | 485 |     |     |     |     | 490 |     |     |     |     | 495 |     |
| Phe | Ala | Ala | Gly | Arg | Val | Ile | Gly | Lys | Gly | Gly | Lys | Thr | Val | Asn | Glu |
|     |     |     | 500 |     |     |     | 505 |     |     |     |     |     | 510 |     |     |
| Leu | Gln | Asn | Leu | Ser | Ser | Ala | Glu | Val | Val | Val | Pro | Arg | Asp | Gln | Thr |
|     |     | 515 |     |     |     |     | 520 |     |     |     |     | 525 |     |     |     |
| Pro | Asp | Glu | Asn | Asp | Gln | Val | Val | Val | Lys | Ile | Thr | Gly | His | Phe | Tyr |



530

535

540

Ala Cys Gln Val Ala Gln Arg Lys Ile Gln Glu Ile Leu Thr Gln Val  
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Arg Arg Lys

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<212> DNA

<213> Homo sapiens

<400> 447

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<212> DNA

<213> Artificial Sequence

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&lt;223&gt; PCR primer

&lt;400&gt; 448

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&lt;210&gt; 449

&lt;211&gt; 579

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 449

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Met | Asn | Lys | Leu | Tyr | Ile | Gly | Asn | Leu | Ser | Glu | Asn | Ala | Ala | Pro | Ser |
|     |     |     | 5   |     |     |     |     |     | 10  |     |     |     |     | 15  |     |

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Asp | Leu | Glu | Ser | Ile | Phe | Lys | Asp | Ala | Lys | Ile | Pro | Val | Ser | Gly | Pro |
|     |     |     | 20  |     |     |     |     | 25  |     |     |     |     | 30  |     |     |

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Phe | Leu | Val | Lys | Thr | Gly | Tyr | Ala | Phe | Val | Asp | Cys | Pro | Asp | Glu | Ser |
|     |     | 35  |     |     |     |     | 40  |     |     |     |     | 45  |     |     |     |

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Trp | Ala | Leu | Lys | Ala | Ile | Glu | Ala | Leu | Ser | Gly | Lys | Ile | Glu | Leu | His |
|     | 50  |     |     |     |     | 55  |     |     |     |     | 60  |     |     |     |     |

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Gly | Lys | Pro | Ile | Glu | Val | Glu | His | Ser | Val | Pro | Lys | Arg | Gln | Arg | Ile |
|     | 65  |     |     |     | 70  |     |     |     |     | 75  |     |     |     |     | 80  |

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Arg | Lys | Leu | Gln | Ile | Arg | Asn | Ile | Pro | Pro | His | Leu | Gln | Trp | Glu | Val |
|     |     |     | 85  |     |     |     |     |     | 90  |     |     |     |     | 95  |     |

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Leu | Asp | Ser | Leu | Leu | Val | Gln | Tyr | Gly | Val | Val | Glu | Ser | Cys | Glu | Gln |
|     |     |     | 100 |     |     |     |     | 105 |     |     |     |     | 110 |     |     |

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Val | Asn | Thr | Asp | Ser | Glu | Thr | Ala | Val | Val | Asn | Val | Thr | Tyr | Ser | Ser |
|     |     |     | 115 |     |     |     | 120 |     |     |     |     |     | 125 |     |     |

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Lys | Asp | Gln | Ala | Arg | Gln | Ala | Leu | Asp | Lys | Leu | Asn | Gly | Phe | Gln | Leu |
|     |     | 130 |     |     |     | 135 |     |     |     |     | 140 |     |     |     |     |

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Glu | Asn | Phe | Thr | Leu | Lys | Val | Ala | Tyr | Ile | Pro | Asp | Glu | Thr | Ala | Ala |
|     | 145 |     |     |     | 150 |     |     |     |     | 155 |     |     |     | 160 |     |

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Gln | Gln | Asn | Pro | Leu | Gln | Gln | Pro | Arg | Gly | Arg | Arg | Gly | Leu | Gly | Gln |
|     |     |     |     | 165 |     |     |     |     | 170 |     |     |     |     | 175 |     |

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Arg | Gly | Ser | Ser | Arg | Gln | Gly | Ser | Pro | Gly | Ser | Val | Ser | Lys | Gln | Lys |
|     |     |     | 180 |     |     |     |     | 185 |     |     |     |     | 190 |     |     |

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Pro | Cys | Asp | Leu | Pro | Leu | Arg | Leu | Leu | Val | Pro | Thr | Gln | Phe | Val | Gly |
|     |     | 195 |     |     |     |     | 200 |     |     |     |     | 205 |     |     |     |

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ala | Ile | Ile | Gly | Lys | Glu | Gly | Ala | Thr | Ile | Arg | Asn | Ile | Thr | Lys | Gln |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|

|         |         |         |         |         |         |         |         |         |         |         |         |         |         |         |         |
|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|---------|
| Thr 225 | Gln 210 | Ser     | Lys     | Ile     | Asp 230 | Val 215 | His     | Arg     | Lys     | Glu 235 | Asn 220 | Ala     | Gly     | Ala     | Ala 240 |
| Glu     | Lys     | Ser     | Ile     | Thr 245 | Ile     | Leu     | Ser     | Thr     | Pro 250 | Glu     | Gly     | Thr     | Ser     | Ala 255 | Ala     |
| Cys     | Lys     | Ser     | Ile 260 | Leu     | Glu     | Ile     | Met     | His 265 | Lys     | Glu     | Ala     | Gln     | Asp 270 | Ile     | Lys     |
| Phe     | Thr     | Glu 275 | Glu     | Ile     | Pro     | Leu     | Lys 280 | Ile     | Leu     | Ala     | His     | Asn 285 | Asn     | Phe     | Val     |
| Gly     | Arg 290 | Leu     | Ile     | Gly     | Lys     | Glu 295 | Gly     | Arg     | Asn     | Leu     | Lys 300 | Lys     | Ile     | Glu     | Gln     |
| Asp 305 | Thr     | Asp     | Thr     | Lys     | Ile 310 | Thr     | Ile     | Ser     | Pro     | Leu 315 | Gln     | Glu     | Leu     | Thr     | Leu 320 |
| Tyr     | Asn     | Pro     | Glu     | Arg 325 | Thr     | Ile     | Thr     | Val     | Lys 330 | Gly     | Asn     | Val     | Glu     | Thr 335 | Cys     |
| Ala     | Lys     | Ala     | Glu 340 | Glu     | Glu     | Ile     | Met     | Lys 345 | Lys     | Ile     | Arg     | Glu     | Ser 350 | Tyr     | Glu     |
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| Asn     | Leu 370 | Asn     | Ala     | Leu     | Gly     | Leu 375 | Phe     | Pro     | Pro     | Thr     | Ser 380 | Gly     | Met     | Pro     | Pro     |
| Pro 385 | Thr     | Ser     | Gly     | Pro     | Pro 390 | Ser     | Ala     | Met     | Thr     | Pro 395 | Pro     | Tyr     | Pro     | Gln     | Phe 400 |
| Glu     | Gln     | Ser     | Glu     | Thr 405 | Glu     | Thr     | Val     | His     | Leu 410 | Phe     | Ile     | Pro     | Ala     | Leu 415 | Ser     |
| Val     | Gly     | Ala     | Ile 420 | Ile     | Gly     | Lys     | Gln     | Gly 425 | Gln     | His     | Ile     | Lys     | Gln 430 | Leu     | Ser     |
| Arg     | Phe     | Ala 435 | Gly     | Ala     | Ser     | Ile     | Lys 440 | Ile     | Ala     | Pro     | Ala     | Glu 445 | Ala     | Pro     | Asp     |
| Ala     | Lys     | Val     | Arg     | Met     | Val     | Ile 455 | Ile     | Thr     | Gly     | Pro     | Pro 460 | Glu     | Ala     | Gln     | Phe     |
| Lys 465 | Ala     | Gln     | Gly     | Arg     | Ile 470 | Tyr     | Gly     | Lys     | Ile     | Lys 475 | Glu     | Glu     | Asn     | Phe     | Val 480 |
| Ser     | Pro     | Lys     | Glu     | Glu 485 | Val     | Lys     | Leu     | Glu     | Ala 490 | His     | Ile     | Arg     | Val     | Pro 495 | Ser     |
| Phe     | Ala     | Ala     | Gly 500 | Arg     | Val     | Ile     | Gly     | Lys 505 | Gly     | Gly     | Lys     | Thr     | Val 510 | Asn     | Glu     |

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